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Expression of pyrrothine *N*-acyltransferase activities in *Saccharothrix algeriensis* NRRL B-24137: new insights into dithiolopyrrolone antibiotic biosynthetic pathway

A.C. Chorin¹, L. Bijeire², M.C. Monje¹, G. Baziard², A. Lebrihi¹ and F. Mathieu¹

1 Université de Toulouse, Laboratoire de Génie Chimique UMR 5503 (CNRS/INPT/UPS), ENSAT INP de Toulouse, 1 Avenue de l'Agrobiopôle, Castanet Tolosan Cedex, France

2 Université de Toulouse, LU 49 Laboratoire de Chimie Pharmaceutique, Faculté des Sciences Pharmaceutiques, Toulouse Cedex, France

Keywords

antibiotic synthases regulation, dithiolopyrrolone biosynthesis, Saccharothrix algeriensis.

Correspondence

Ahmed Lebrihi, Université de Toulouse, Laboratoire de Génie Chimique UMR 5503 (CNRS/INPT/UPS), ENSAT INP de Toulouse, 1 Avenue de l'Agrobiopôle, B.P. 32607, F 31326 Castanet Tolosan Cedex 1, France. E mail: lebrihi@ensat.fr

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Abstract

Aims: The hypothetical dithiolopyrrolone biosynthetic pathway includes a final step of pyrrothine nucleus acylation. The presence of an enzymatic activity catalysing this reaction was investigated in *Saccharothrix algeriensis* NRRL B 24137. To understand the effect exerted by organic acids on the level of dithiolopyrrolone production, their influence on enzymatic expression was studied.

Methods and Results: The transfer of acetyl CoA or benzoyl CoA on pyrro thine was assayed in the cell free extract of *Sa. algeriensis* NRRL B 24137. This study reports the presence of an enzymatic activity catalysing this reaction that was identified as either pyrrothine *N* acetyltransferase or *N* benzoyltransferase. The stimulation of benzoyl pyrrothine (BEP) production by addition of benzoic acid at 1.25 mmol 1⁻¹ into the culture medium was demonstrated, and results showed that under the same conditions of growth, pyrrothine *N* benzoyltransferase specific activity was doubled.

Conclusions: This study shows that BEP production is enhanced in the presence of benzoic acid partly because of an induction of pyrrothine N benzoyltransfer ase.

Significance and Impact of the Study: The antitumor and antibiotic properties of dithiolopyrrolones are related to their variable acyl groups. New insights into regulation of biosynthetic pathway, especially the step of pyrrothine acylation, could lead after further studies to yield improvement and to selective production of dithiolopyrrolones with new biological activities.

Introduction

Saccharothrix algeriensis NRRL B 24137 is a filamentous bacterium that produces bioactive metabolites belonging to the dithiolopyrrolone class of antibiotics (Lamari *et al.* 2002a,b; Zitouni *et al.* 2005). Dithiolopyrrolones possess a common pyrrolinonodithiole nucleus linked to two vari able groups R1 and R2 (Fig. 1a). Saccharothrix algeriensis NRRL B 24137 produces at least six pyrrothine deriva tives characterized by their different N acyl groups R2: thiolutin (acetyl pyrrothine), senecioyl pyrrothine, tigloyl pyrrothine, isobutyryl pyrrothine, butanoyl pyrrothine and benzoyl pyrrothine (BEP) (Lamari *et al.* 2002a,b) (Fig. 1b). Pyrrothine derivatives differ from holothin derivatives by the presence of a methyl radical R1 linked to the N_4 cyclic nitrogen.

Many bacteria have been identified as dithiolopyrro lones producers among the genera *Streptomyces* (Celmer *et al.* 1952; Bhate *et al.* 1960), *Xenorhabdus* (Li *et al.* 1995) and *Alteromonas* (Shiozawa *et al.* 1993); however, *Sa. algeriensis* NRRL B 24137 is the first producer identi fied within the *Saccharothrix* genus. In addition, naturally occurring and other dithiolopyrrolones can also be pro duced by multiple step chemical synthesis (Büchi and Lukas 1964; Hagio and Yoneda 1974; Ellis *et al.* 1977; Hjelmgaard *et al.* 2007; Li *et al.* 2007).



Figure 1 (a) General structure of dithiolopyrrolones: the pyrrolinonodithiole nucleus and its radicals R1 and R2. (b) Structure of dithiolopyrrolones produced by Saccharothrix algeriensis NRRL B 24137.

Dithiolopyrrolones exhibit broad spectrum antibiotic activity against a variety of Gram negative and Gram positive bacteria, fungi (Lamari et al. 2002a) and even protozoa and insects (Mc Inerney et al. 1991). They also inhibit allergy (Stahl et al. 1988) and platelet aggregation (Ninomiya et al. 1980). Moreover, dithiolopyrrolones generated new interests in the 1990s after the discovery of their antitumour properties. Thiolutin suppresses tumour cell induced angiogenesis (Minamigushi et al. 2001), and a variety of dithiolopyrrolones show strong cytotoxic activities against human cancer cell lines, especially breast, colon and cervical ones (Webster et al. 2000). Guo et al. (2008) have recently described some new chemically syn thesized dithiolopyrrolones that promote production of white blood cells. They could be useful in the prevention and treatment of microbial infections such as HIV infec tions and for the treatment of blood disorders.

Biological activity of dithiolopyrrolones is strongly influenced by the nature of variable groups (Oliva et al. 2001; Chen et al. 2006; Li et al. 2007; Guo et al. 2008). As a consequence, it is important to have the means to influence the type of dithiolopyrrolone produced. Unfor tunately, little is known about the dithiolopyrrolone biosynthetic pathway to date. A single putative pathway was described by Furumai et al. (1982) (Fig. 2). L cystine has been identified as the potential precursor of pyrro thine nucleus. Amide bond formation between pyrro thine and an activated organic acid would lead to dithiolopyrrolone synthesis. The presence of one enzy matic activity that catalyses amide bond formation between the holothine nucleus (deacetyl holomycin) and acetyl CoA was confirmed in cell free extracts of Strepto myces clavuligerus mutants, which are producers of holo mycin (De la Fuente et al. 2002), but the enzyme responsible for this activity has not been purified yet. Additionally, the influence of medium composition has provided some information on the biosynthetic pathway in *Sa. algeriensis* NRRL B 24137. Addition of supposed precursors into the culture medium, such as organic and amino acids, led to modifications in dithiolopyrrolone production levels (Bouras *et al.* 2006a,b, 2007) and to precursor directed biosynthesis of new dithiolopyrrolone analogues (Bouras *et al.* 2008). However, little is known about enzymes involved in the biosynthesis of dithiolo pyrrolones by this strain.

To get a better understanding of dithiolopyrrolone biosynthetic pathway in *Sa. algeriensis* NRRL B 24137, the enzymatic reaction of pyrrothine nucleus acylation (pyrrothine *N* acyltransferase activity) was investigated. For the first time, this study looks into the mechanism involved in the precursor directed biosynthesis of dithiolopyrrolones by *Sa. algeriensis* NRRL B 24137. It leads to new insights into biosynthetic pathway regulation and thus could result in production yield improvement and to selective production of new and specific dithiolo pyrrolones.

Materials and methods

Producing strain

Saccharothrix algeriensis NRRL B 24137 (=DSM 44581) was used for this study. A stock of spores was prepared to maintain the strain. Saccharothrix algeriensis NRRL B 24137 was grown on International Streptomyces Project 2 (ISP2) agar plates for 7 days at 30°C. Spores were suspended in 0.1% Tween 80 (Fisher, Waltham, MA), harvested and stored in 25% glycerol (Fisher) at -24°C.



Figure 2 Dithiolopyrrolone biosynthetic pathway (according to Furumai *et al.* 1982).

Culture media

ISP2 was composed (per litre of distilled water) of 10 g D (+) glucose (Acros, Geel, Belgium), 10 g malt extract (Difco), 4 g yeast extract (Difco) and 18 g agar (Difco). The pH was adjusted to 7 with 2 N NaOH (Sigma) before autoclaving for 20 min at 121° C.

Semi synthetic (SS) medium (SSM) developed in our laboratory was used for growth and antibiotic production (Bouras *et al.* 2006a) with 15 g l⁻¹ of D (+) glucose (Acros). Organic acids were autoclaved separately and added aseptically to the culture medium, at the required concentration, before inoculation. The effect of organic acids on the time course of antibiotic production and enzymatic expression was investigated with 1.25 mmol l⁻¹ of benzoic acid (Acros) or 5 mmol l⁻¹ of acetic acid (Sigma). The influence of organic acid concentration on maximal enzymatic expression was studied by the addition of benzoic acid at concentrations of 0, 0.5, 2.5 and 5 mmol l⁻¹ and by the addition of acetic acid at concentrations of 0, 2.5, 5 and 7.5 mmol l⁻¹.

Culture conditions

ISP2 agar plates were inoculated with 50 μ l of spore stock, i.e. 2 × 10⁷ CFU, and incubated for 7 days at 30°C for spore production. Precultures were inoculated with spores harvested from one ISP2 agar plate. They were grown in 250 ml Erlenmeyer flasks containing 50 ml of SSM and incubated during 48 h. Cultures were grown in 500 ml Erlenmeyer flasks containing 100 ml of SSM, which was, in some cases, supplemented with organic acids. They were inoculated with 5 ml of homogenized preculture. All liquid cultures were incubated at 30°C and 240 rev min¹ on a rotary shaker (New Brunswick Scientific Company, New Brunswick, NJ, USA).

Dry cell weight measurement

Culture broth samples of 4 ml were taken every 12 h of fermentation and centrifuged at 16 000 g for 10 min in preweighed Eppendorf tubes (centrifuge Biofuge pico; Heraeus Instruments, Hanau, Germany). Dry cell weights (DCWs) were determined as described by Bouras *et al.* (2006a) and expressed in g 1⁻¹.

Dithiolopyrrolone extraction

A volume of 3.2 ml of supernatant fluid was extracted twice with 1.6 ml of dichloromethane. Organic phases containing antibiotics were pooled and dried under vacuum in a speed vac (Genevac, Ish, UK) at a tempera ture maintained under 40°C. Dry extracts were dissolved in 0.8 ml of methanol (HPLC grade; Fisher) for dith iolopyrrolone quantification by high performance liquid chromatography (HPLC).

Pyrrothine synthesis

PMBS

6 amino 4 methyl [1,2]dithiolo[4,3 b]pyrrol 5 one hydro chloride, also called pyrrothine, was synthesized by multi ple step chemical synthesis using the protocol of Hjelmgaard *et al.* (2007) with slight modifications as described in Fig. 3. Especially, the methyl group on cyclic nitrogen was introduced during the step (b) by substitut ing *p* methoxybenzylamine by methylamine. Finally, the amide moiety of the trifluoroacetyl pyrrothine (5) was hydrolysed (f) in methanol in the presence of concen trated HCl to afford the pyrrothine hydrochloride (6) that

PMBS



was recovered by filtration. The ¹H NMR spectrum, obtained on a Bruker DPX 300 MHz spectrometer, con firmed that the precipitate is pyrrothine hydrochloride: $\delta_{\rm H}$ (DMSO, d₆): 7·41 (1H, s, C=CH) 3·9 5·3 (1H, br s, NH₃⁺) 3·27 (3H, s, CH₃).

Cell free extract preparation

Saccharothrix algeriensis NRRL B 24137 was grown in SSM or SSM supplemented with organic acids as described in the section 'Culture conditions'. The biomass was recovered by centrifugation of culture broth at 5000 g for 15 min (4K15; Sigma), then washed twice with physiological water (0.9% NaCl) and once with lysis buffer (Tris HCl 50 mmol 1¹, pH 8). Wet cells were finally recovered by filtration on 0.2 μ m membrane filters (Advantec, Dublin, Ireland). Wet cells (0.6 g) were re suspended in 1 ml of lysis buffer and transferred to a Fast Protein Blue tube (MP Biomedicals, Irvine, CA, USA). Two disruption cycles (30 s, 5 m s⁻¹) were carried out in a Fast Prep disruptor (MP Biomedicals). The lys ing matrix was discarded, and then the sample was centri fuged at 10 000 g for 30 min (centrifuge 1 15K; Sigma) to remove the cell debris. The supernatant constituting the soluble cell free extract of Sa. algeriensis NRRL B 24137 was used immediately for the assay of pyrrothine N acyltransferase activity. A sample was frozen at -80°C for further protein assays. Samples were always kept in ice and centrifugations were carried out at 4°C.

Pyrrothine N acyltransferase assay

Pyrrothine *N* acyltransferase activity was assayed in a final volume of 100 μ l. The reaction mixture contained 80 μ l of cell free extract in Tris HCl buffer (50 mmol l⁻¹, pH 8), 10 μ l of acetyl or benzoyl coenzyme A at 5 mmol l⁻¹ in bidistilled water and 10 μ l of pyrrothine hydrochoride at 2.5 mmol l⁻¹ in methanol. The reaction mixture was incubated at 30°C for 5 and 10 min. The reaction was stopped by adding fresh 2.5% w/v trichloroacetic acid (Fisher). The cell free extract was properly diluted with Tris HCl buffer (50 mmol l⁻¹, pH 8) before assay to observe a product formation linear in time within 10 min.

Enzymatic activity was identified as either acetyltrans ferase or benzoyltransferase activity according to the acyl group donor used during the assay, i.e. acetyl CoA and benzoyl CoA, respectively. A unit of enzyme is defined as the enzyme activity producing 1 μ mol of thiolutin (acetyl pyrrothine) or BEP per minute. Specific enzymatic activity was expressed in μ U mg⁻¹ of protein.

Enzymatic activity was also expressed in mU g¹ of DCW. A value of 162 mg of proteins per gram of dry

cells was used to achieve conversion from U mg⁻¹ of proteins to U g⁻¹ of DCW. Protein content in dry cells was determined as follow. DCW and intracellular protein concentration were determined every 24 h, for 7 days, in a culture of *Sa. algeriensis* NRRL B 24137 on SSM. The plot of proteins in mg l⁻¹ vs DCW in g l⁻¹ indicated a linear correlation between the two parameters with a slope of 162 mg of proteins per gram of DCW ($R^2 = 0.91$).

HPLC analysis for dithiolopyrrolone quantification

Dithiolopyrrolones were quantified by HPLC (Bio tek Instruments, Milan, Italy). The analytical column was ProntoSIL 120 5 C_{18} SH, 150 × 4.6 mm (Bishoff Chroma tography, Leonberg, Germany) fitted with a guard column of 10 × 4 mm, and detection was achieved with a diode array detector (UV vis 545 V; Bio tek instruments).

For detection and quantification of thiolutin and BEP produced in the culture medium, analyses were per formed as described by Bouras *et al.* (2006a) with an injection volume of 80 μ l.

For detection and quantification of dithiolopyrrolones formed during enzymatic assay, analyses were performed under the following chromatographic conditions. Samples were analysed by linear gradient elution using a mixture of methanol/bidistilled water (solvent A/solvent B) as mobile phase and a flow rate of 0.8 ml min¹. Column temperature was maintained at 30°C and injection volume was 40 μ l. UV detection of antibiotics was carried out at 390 nm. For the acetyltransferase assay, elution was carried out with a linear gradient from 30% A to 47% A in 17 min. Pyrrothine and thiolutin retention times (R_t) were 7 and 12.3 min, respectively. For the benzoyltrans ferase assay, the mobile phase composition was 30% A, reached 45% A in 15 min, 100% A in 35 min and was finally kept at this value for 2 min. Pyrrothine and BEP retention times (R_t) were 8 and 29 min, respectively.

Both antibiotics were quantified using a thiolutin standard calibration curve. Indeed, the molar absorbtivity values at 390 nm of pyrrothine derivatives are close. e^{390} is in the range of 8317 9333 mol⁻¹ l cm⁻¹ as described by Lamari *et al.* (2002b).

The BEP chemical structure was confirmed by compar ison of the UV spectrum obtained with those mentioned by Bouras *et al.* (2008). The UV spectrum of BEP yielded λ_{max} in nm (relative absorbance): 231 (1), 309 (0.45), 401 (0.68).

Replication of experiments and statistical analysis

Data of growth and dithiolopyrrolone production repre sent the average of triplicate flasks, and error bars denote standard errors. Values of acyltransferase specific activities at 24 and 48 h on SSM, SSM with benzoic acid at 1.25 mmol l^{-1} and SSM with acetic acid at 5 mmol l $^{-1}$ are the averages of four independent experiments, and error bars denote standard errors. Other pyrrothine N acyltransferase specific activities are single values, and error bars denote a standard error of 15%, estimated from previous replications.

Statistical analyses were achieved using KALEIDA GRAPH 4.0 software (Synergy Software, Reading, PA, USA).

Results

Involvement of pyrrothine N acyltransferase activities in thiolutin and BEP production

N acyltransferase enzymatic activity was assayed in the cell free extract of *Sa. algeriensis* NRRL B 24137 grown on SSM. A formation of thiolutin or BEP, linear in time within 10 min, was monitored by HPLC when acetyl CoA and benzoyl CoA were respectively used as acyl group donors (Fig. 4). No thiolutin or BEP was observed in the absence of one of the components of the reaction mix ture, i.e., acyl CoA, cell free extract or pyrrothine. Hence, it can be concluded that both pyrrothine *N* acetyltransfer ase and pyrrothine *N* benzoyltransferase enzymatic activi ties are present in the cell free extract of *Sa. algeriensis* NRRL B 24137.

Biomass, thiolutin and BEP productions on SSM

DCW and dithiolopyrrolone concentrations were quanti fied during growth on SSM (Fig. 5a). Time courses of specific growth rate, thiolutin specific production and BEP specific production on SSM are shown in Fig. 6a.

Two distinct periods of growth were observed. Each growth period is characterized by an increase in the spe

cific growth rate followed by a decrease in the same. Between 0 and 48 h, specific growth rate reached the maximal value of 0.096 h⁻¹ at 12 h and then decreased up to 0.010 h⁻¹ at 48 h. At the end of this first period of growth, a DCW of 1.2 g l⁻¹ was observed. Between 60 and 96 h, the specific growth rate increased again to the maximal value of 0.022 h⁻¹ at 60 h and then decreased to zero at 96 h. The maximal specific growth rate during this second period was 4.4 times lower than the earlier one. The maximal level of DCW, 2.5 g l⁻¹, was attained at 96 h. Biomass finally dropped after 108 h probably because of cell lysis.

Thiolutin and BEP production started at 24 and 36 h of culture, respectively. At 72 h of culture, the maximal level of thiolutin, 25.6 mg l⁻¹, was observed. Then, the thiolutin concentration in the culture medium decreased. A BEP production of 0.15 mg l⁻¹ was reached as early as 48 h, i.e. 80% of the maximal BEP concentration, 0.18 mg l⁻¹, observed only at 144 h. Maximal specific productions of thiolutin and BEP were reached after 48 h of culture at the end of the first growth period, when the specific growth rate was at its lowest level, 0.010 h⁻¹. Maximal specific productions of 20 and 0.12 mg g⁻¹ were respectively observed for thio lutin and the BEP. The maximal specific production rate of thiolutin, 0.70 mg g 1 h 1, was reached at 24 h and the maximal specific production rate of BEP, 0.006 mg g⁻¹ h⁻¹, was observed at 36 h when the spe cific growth rate was reduced to 26.5 and 12% of its maximal level, respectively.

Addition of benzoic acid and acetic acid to the culture medium

Time courses of growth, thiolutin production and BEP production on SSM completed with benzoic acid at









1·25 mmol l¹ are shown in Fig. 5b. Time courses of spe cific growth rate, thiolutin specific production and BEP specific production on SSM supplemented with benzoic acid at 1·25 mmol l¹ are shown in Fig. 6b. A maximal BEP production of 8·47 mg l¹ was observed at 72 h. This is 56 times more than on SSM. We observed a BEP maxi mal specific production of 4·77 mg g¹ at 60 h, i.e. a value 39 times higher than without benzoic acid in cul ture medium. Additionally, the maximal specific produc tion rate of BEP, 0·15 mg g¹ h¹, at 36 h, was 25 times higher than the one on SSM. By contrast, thiolutin pro duction and growth were not significantly modified by benzoic acid addition.



Figure 6 Time course of specific growth rate (μ, \diamond) , specific production of thiolutin (\bigcirc) and specific production of benzoyl pyrrothine (\bigtriangleup) during cultures of *Saccharothrix algeriensis* NRRL B 24137 on (a) SS medium (SSM), (b) SSM supplemented with benzoic acid at 1.25 mmol l⁻¹, (c) SSM supplemented with acetic acid at 5 mmol l⁻¹. Specific growth rate is given in h⁻¹, specific productions in mg g⁻¹ of dry cell weight and time in hours.

DCW and dithiolopyrrolone concentrations were also quantified in a *Sa. algeriensis* NRRL B 24137 culture on SSM completed with acetic acid at 5 mmol 1⁻¹ (Fig. 5c). Specific growth rate, thiolutin specific production and BEP specific production are shown in Fig. 6c. A maximal thiolutin concentration of 9.68 mg l⁻¹ was observed at 72 h, which is 2.6 times less than on SSM. Data also showed that thiolutin specific production reached a maxi mal value of 6.53 mg g⁻¹ at 48 h, i.e. three times lower than that on SSM. The maximal specific production rate of thiolutin was also reduced by acetic acid addition. The value of 0.23 mg g⁻¹ h⁻¹, three times lower than the one on SSM, was reached at 24 h. On the contrary, acetic acid addition did not significantly modify BEP production and growth.

Expression of pyrrothine N acyltransferase activities on SSM

Time courses of acetyltransferase and benzoyltransferase specific activities during growth on SSM are shown in Fig. 7a. Acetyltransferase specific activity peaked at 24 h and the peak value was 18 823 μ U mg⁻¹ of protein. Then, enzymatic specific activity decreased sharply. At 48 h, the expression was already seven times lower.

Benzoyltransferase activity was detected from 24 h and reached its maximal value of 9450 μ U mg⁻¹ of



protein after 48 h of culture, while acetyltransferase spe cific activity was already reduced to 40% of its maximal level. Then it decreased down to 1760 μ U mg⁻¹ at 84 h. At 96 h of culture a new increase in specific activity was observed until the maximal value of 3690 μ U mg⁻¹ of protein was reached. Then activity decreased down to 1800 μ U mg⁻¹ at 144 h and remained stable until 168 h of culture.

From both profiles comparisons, it is particularly noteworthy that acetyltransferase and benzoyltransferase specific activities did not peak at the same time and that the ratio of both enzymatic specific activities was not constant throughout culture on SSM.

Acetyltransferase specific activity was also compared to the specific production rate of thiolutin (Fig. 8a). The time profile of acetyltransferase specific activity is tightly related to the time profile of thiolutin specific production rate. Both acetyltransferase specific activity and thiolutin specific production rate peaked at 24 h and decreased until 72 h.



Figure 7 Time course of pyrrothine *N* acetyltransferase (\blacksquare) and pyrrothine *N* benzoyltransferase (\Box) specific activities during cultures of *Saccharothrix algeriensis* NRRL B 24137 on (a) SS medium (SSM), (b) SSM supplemented with benzoic acid at 1-25 mmol l⁻¹, (c) SSM supplemented with acetic acid at 5 mmol l⁻¹. Specific activities are given in μ U mg⁻¹ of proteins and time in hours.

Figure 8 Time course of pyrrothine *N* acyltransferase specific activity (\bullet) and rate of average dithiolopyrrolone specific production (\bigcirc) on SS medium. (a) Comparison of acetyltransferase specific activity and rate of average thiolutin specific production. (b) Comparison of benzoyltransferase specific activity and rate of average benzoyl pyrrothine specific production. Pyrrothine *N* acyltransferase specific activities are given in mU g⁻¹ of dry cell weight (DCW) and rates of average dithiolopyrrolone specific production in mg h⁻¹ g⁻¹ of DCW. Time is given in hours.

Similarly, BEP specific production rate is tightly related to the level of benzoyltransferase specific activity (Fig. 8b). The peak of BEP specific production rate is associated with an increase in benzoyltransferase specific activity.

Expression of pyrrothine N acyltransferase activities on SSM with organic acids

Acetyltransferase and benzoyltransferase specific activities were quantified during growth on SSM supplemented with benzoic acid at 1.25 mmol l^{-1} (Fig. 7b) or SSM supplemented with acetic acid at 5 mmol 1^{-1} (Fig. 7c).

During growth on SSM with benzoic acid, benzoyl transferase specific activities were significantly higher than on SSM but the shape of the curve was not modified. Benzoyltransferase specific activity reached a maximal value of 20 881 μ U mg⁻¹ of proteins at 48 h, i.e. 2·2 times higher than without benzoic acid. The second peak of benzoyltransferase specific activity observed at 96 h, like that on SSM, had a value of 15 920 μ U mg⁻¹ of pro teins, i.e. 1·8 times higher than without benzoic acid. On the contrary, acetyltransferase specific activities were not significantly modified by benzoic addition to culture medium.

Additionally, benzoyltransferase specific activity was also determined in cell free extracts obtained after 48 h of growth, on SSM with different concentrations of benzoic acid (Fig. 9a). Benzoyltransferase specific activity was two times higher with benzoic acid added to the culture medium than without, regardless of the benzoic acid concentration, which range from 0.5 to 2.5 mmol 1⁻¹.

The time course of acetyltransferase specific activity was not significantly modified by acetic acid addition to SSM. Acetyltransferase specific activity reached a maximal value of 16 391 μ U mg⁻¹ of protein at 24 h. Similarly, acetic acid had no effect on the time course of benzoyl

transferase specific activity. The maximal benzoyltransfer as e specific activity, 9840 μ U mg ¹ of protein, was reached at 48 h.

Acetyltransferase specific activity was also determined in cell free extracts obtained after 24 h of growth on SSM with different concentrations of acetic acid (Fig. 9b). Acetic acid had no effect on acetyltransferase specific activity regardless of the concentration of acetic acid added to the medium in the range from 2.5 up to 7.5 mmol l¹.

Discussion

Saccharothrix algeriensis NRRL B 24137 has the ability to produce a wide range of dithiolopyrrolones with different radicals (R2s) depending on the precursors added to the culture medium (Lamari et al. 2002a,b; Bouras et al. 2006a, 2008). To have such ability to produce many dithiolopyrrolones with broad structural diversity, the micro organism must possess not only precursors but also a suitable enzymatic system with sufficient flexibility to attach a variety of R2 radicals on N7. Data presented above provide evidence for the presence in the cell free extract of Sa. algeriensis NRRL B 24137 of an enzymatic activity that catalyses acylation of pyrrothine nucleus on N7 using acyl CoA as acyl group donors. Especially, acetyl CoA and benzoyl CoA are respectively substrates of a pyrrothine N acetyltransferase and N benzoyltransferase enzymatic activity. These results suggest that the enzy matic reaction of pyrrothine acylation takes part in the dithiolopyrrolone biosynthetic pathway in Sa. algeriensis NRRL B 24137, which is able to use acyl CoA with very different structures (acetyl CoA and benzoyl CoA), as substrates to produce corresponding dithiolopyrrolones (thiolutin and BEP). The presence of a similar enzymatic activity, called holomycin synthase, catalysing the amide bond formation between holothin (deacetyl holomycin)



Figure 9 Influence of acid concentration on *N* pyrrothine acyltransferase activities. (a) Influence of benzoic acid concentration in SS medium (SSM) on pyrrothine *N* benzoyltransferase specific activity at 48 h. (b) Influence of acetic acid concentration in SSM on pyrrothine *N* acetyltransferase specific activity at 24 h. Specific enzymatic activities are given in μ U mg⁻¹ of proteins and acid concentrations in mmol l⁻¹.

and acetyl CoA was already observed in the cell free extracts of *S. clavuligerus* mutants, which are producers of holomycin (De la Fuente *et al.* 2002). However, the enzyme responsible for this activity has not been purified yet. In contrast, the existence of N benzoyltransferase activity involved in dithiolopyrrolone synthesis has never been reported before.

To understand the regulatory mechanisms exerted on these new enzymatic activities, the time courses of pyrro thine N acetyltransferase and pyrrothine N benzoyltrans ferase specific activities were determined in the cell free extract of *Sa. algeriensis* NRRL B 24137 and compared to the time courses of formation of thiolutin, BEP and growth.

Data shown in this study indicate that the dith iolopyrrolone production phase is partly growth associ ated during culture of Sa. algeriensis NRRL B 24137 on SSM and occurs during a decrease in the specific growth rate. These results contrast with the traditional idea that the antibiotic production phase, called the idiophase, is separated from the growth phase, called the trophophase. However, several studies have shown that culture condi tions supporting slow growth rates, e.g. nutritional limita tions, can elicit an antibiotic production partly associated with growth (Martin and Demain 1980; McDermott et al. 1993; Fazeli et al. 1995; Untrau Taghian et al. 1995). Besides, studies on antibiotic production using chemostat cultures demonstrated that production can occur in asso ciation with growth although antibiotic production level is higher at low dilution rates (Vu Trong and Gray 1981; Lebrihi et al. 1988; McIntyre et al. 1996; Pamboukian and Facciotti 2004).

In addition, a noticeable decrease in thiolutin concen tration in the culture medium is observed after 72 h of growth on SSM. It is associated with a decrease in antimi crobial activities against Muccor ramaniannus and Bacillus subtilis (L. Lamari, unpublished data). Thiolutin could undergo a physical or a chemical transformation (Yoshioka and Stella 2007). This compound contains in particular a disulfide bond, which could be chemically broken with a relative ease (Parker and Kharasch 1959; Kice 1968). Thiolutin could also be converted enzymati cally as already described in the literature for other antibi otics. Enzymatic conversion could be either extracellular or intracellular after reassimilation of thiolutin by the cells (Argoudelis and Mason 1969; Perlman and Sebek 1971). However, no mechanism of thiolutin transforma tion has been identified yet.

Pyrrothine N acyltransferase activity was detected throughout the culture but was only overexpressed during a very short time that coincides with the peak of anti biotic productivity (i.e. specific production rate). Such expression profiles have been previously observed for dif ferent antibiotic synthases (Ortmann *et al.* 1974; Nimi *et al.*1981; Brana *et al.* 1985). It is an illustration, at the biochemical level, that antibiotic synthesis is very tightly regulated as it has been reported in the study of antibiotic gene clusters regulation in *Streptomyces* sp. (Cundliffe 2006; Rokem *et al.* 2007).

Saccharothrix algeriensis NRRL B 24137 showed a valuable capability to produce new dithiolopyrrolone analogues when adequate precursors were added into the culture medium (Bouras *et al.* 2008). Organic acid addi tion promoted the production of new dithiolopyrrolones possessing a R2 radical either identical to or different from organic acid. Specifically, benzoic acid or cinnamic acid addition led to BEP production and valeric acid addition resulted in valeryl pyrrothine production (Bouras *et al.* 2008).

In our study, we showed that BEP was produced at a very low level on SSM and we confirmed that BEP pro duction was strongly increased by benzoic acid addition. Additionally, analysis of pyrrothine N benzoyltransferase specific activity showed that specific activity was markedly enhanced during growth in the presence of benzoic acid. Benzoic acid had no activator effect on benzoyltransferase specific activity when added in the benzoyltransferase assay (data not shown). These results suggest that stimu lation of BEP production by benzoic acid is not only because of an additional precursor supply but also includes the induction of a biosynthetic enzymatic activity, i.e. pyrrothine N benzoyltransferase, responsible for the transfer of benzoyl CoA to pyrrothine.

Other antibiotic precursors that stimulate antibiotic production by inducing biosynthetic enzymes are known. A typical example is the stimulation of ergot alkaloid production by tryptophan. Krupinski *et al.* (1976) showed that tryptophan enhanced the alkaloid produc tion of ergot by inducing dimethylallyltryptophan synthe tase, the first enzyme in the ergot alkaloid biosynthetic pathway.

In addition, earlier studies provided evidence that methi onine, a precursor of cephalosporin C in *Cephalosporium acremonium*, induces enzymes of the cephalosporin C biosynthetic pathway such as δ (L α aminoadipyl) L cys teinyl D valine synthetase, isopenicillin N synthase and deacetoxycephalosporin C synthase (Demain and Zhang 1998). Regulation seems to occur at the transcriptional level because the transcription of genes encoding the above enzymes was strongly increased by culture on methionine (Demain and Zhang 1998). Lastly, it has been shown that some enzymes involved in pathways supplying antibiotic precursors, especially amino acid catabolism pathways, are induced by their substrate. Rius *et al.* (1996) observed that lysine induced the L Lysine ε amino transferase, an enzyme involved in the conversion of L lysine to L α aminoadipic acid, a direct precursor of cephalosporins in *S. clavuligerus*. Besides, valine deshydrogenase, the first enzyme of valine catabolism that provides *N* butyrate, propionate and methyl malonate for macrolide biosynthesis is induced by valine in *Strepto myces aureofaciens*, *Streptomyces fradiae* and *Streptomyces avermitilis* (Nguyen *et al.* 1995a,b).

By contrast, acetic acid addition to the culture medium depressed thiolutin production despite the fact that it is a potential precursor for thiolutin synthesis. Our assays of the pyrrothine N acetyltransferase activity showed that acetic acid had no effect on its level of activity, support ing the idea that the decrease in thiolutin production was not because of repression of this enzymatic activity. Acetic acid had no inhibitor effect on acetyltransferase specific activity when added in the acetyltransferase assay (data not shown). As a consequence, the negative effect of acetate addition on thiolutin specific production remains to be elucidated but some hypothetical mecha nisms can be put forth. Possible explanations include an inhibition of the acetyltransferase activity by a change in intracellular pH, a negative effect of acetate on another enzyme of the biosynthetic pathway or on enzymes involved in a pathway supplying antibiotic precursors. Modifications of transport systems or redirection of thio lutin precursors to primary metabolism might also account for a decrease in thiolutin production in the presence of acetate but it is unlikely because growth was not affected by acetate addition.

Lastly, some of the results presented in this study sup port the idea that the transfer reactions of acetyl CoA and benzoyl CoA on pyrrothine are not catalysed by the same enzyme. The existence of a unique enzyme could not account for the variability in the ratio between both specific activities throughout the culture. Additionally, this idea is strengthened by the finding that the benzoyl transferase activity level is specifically increased during culture with benzoic acid, whereas acetyltransferase activity is not modified.

The purification of enzymes catalysing the transfer reaction of acetyl CoA and benzoyl CoA to pyrrothine is in progress in our laboratory in order to confirm the hypothesis that two different enzymes are involved. Addi tionally, the amino acid sequence determination of enzyme(s) would be an essential starting point for gene identification and further studies on biosynthetic pathway regulation. Further studies should also be carried out on *Sa. algeriensis* NRRL B 24137 nutritional requirements and on its behaviour in a controlled environment within a bioreactor to better understand the conditions of anti biotic onset.

This study provides evidence for the presence of a pyrrothine N acyltransferase activity in the cell free

extract of *Sa. algeriensis* NRRL B 24137 involved in the dithiolopyrrolone biosynthetic pathway. Our work also suggests that enhancement of BEP production by *Sa. alge riensis* NRRL B 24137 in the presence of benzoic acid is partly because of an increase in pyrrothine *N* benzoyl transferase specific activity. As a consequence, this study provides new insights into dithiolopyrrolone synthesis, especially into its regulation and also gives valuable experimental tools to further our understanding of the dithiolopyrrolone biosynthetic pathway.

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