

# An intracellular esterase from *Bacillus cereus* catalyzing hydrolysis of 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol

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**Abstract** An esterase from *Bacillus cereus* that hydrolyzes 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol was purified to homogeneity. After purification, the molecular mass of the esterase was determined as 43 kDa by SDS-PAGE, and estimated as 45 kDa using gel filtration, suggesting that the enzyme is a monomer. The optimum pH and temperature for activity of the enzyme were 7.0 and 40°C respectively. The N-terminal sequence was determined. Whole cells of this strain were applied to the resolution of 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol.

**Keywords** *Bacillus cereus* · Esterase · 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol · Propranolol

## Introduction

$\beta$ -Adrenergic blocking agents such as (S)-propranolol are used mainly to treat angina pectoris, hypertension and other cardiac diseases (Ariens 1984); the desirable therapeutic activities reside mainly in the S-enantiomers of these compounds (Howe and Shanks 1968). Approaches to the preparation of (S)-propranolol can be based on chemical (Klunder et al. 1986; Sharpless et al. 1992; Hiroaki et al. 1993; Veloo and Koomen 1993; Chang and Sharpless 1996;

Takuya et al. 2001) or chemoenzymatic (Noritada and Nobuo 1985; Fuganti et al. 1986; Bianchi et al. 1988; Yoshiyasu et al. 1988; Bevinakatti and Banerji 1991; Chiou et al. 1997; Pchelka et al. 2000; Subhash et al. 2000) methods.

1-Chloro-3-(1-naphthoxy)-2-propanol and its *O*-acetate derivative are key chiral intermediates in the chemoenzymatic methods for the synthesis of propranolol. Resolution of the intermediates can be achieved in aqueous phase or organic solvents. Lipases of different origins, e.g., lipase from *Pseudomonas* species (Hsu et al. 1990), *Pseudomonas cepacia* (Bevinakatti and Banerji 1991), *Mucor miehei* (Ader and Schneider 1992), *Candida antarctica* (Kaimal et al. 1992), *Candida cylindracea*, *Mucor javanicus*, *Trichosporon* sp., *Bacillus* sp. (Kapoor et al. 2003) and *Pseudomonas fluorescens* (Di Nunno et al. 2000) have been successfully resolved. In this work, a *Bacillus cereus* strain expressing an intracellular esterase activity was identified using 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol as the prochiral substrate. Whole cells of the strain were used for kinetic resolution of the substrate (Fig. 1). The esterase was also purified and characterized.

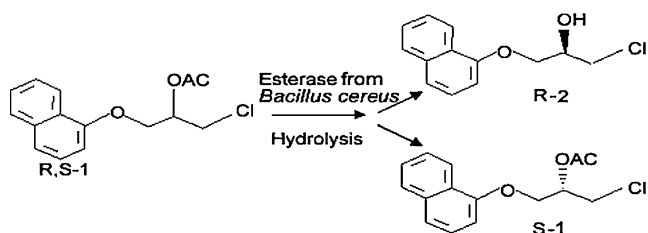
## Materials and methods

### Materials

The two enantiomers of propranolol and para-nitrophenyl fatty acid esters were purchased from Sigma (St. Louis, MO), 1-chloro-3-(1-naphthoxy)-2-propanol and 1-chloro-3-(1-naphthoxy)-2-acetoxy propanol were synthesized and purified in our laboratory according to described methods (Bevinakatti and Banerji 1991). The compound structures were confirmed by IR and NMR spectroscopy.

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**Fig. 1** Esterase-catalyzed resolution of racemic 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol. *S-1* 1-Chloro-3-(1-naphthoxy)-2-acetoxypropanol, *R-2* 1-chloro-3-(1-naphthoxy)-2-propanol

### Microorganism

Strain WL 24 was isolated from soil samples from the suburbs of Beijing city employing a screening agar medium containing 1.0% olive oil, 2.0% yeast extract, 0.1%  $K_2HPO_4$ , 0.05%  $MgSO_4$ , 0.05%  $CaCl_2$ , 0.5% Tween 80 and 1.5% agar. The strain was identified as *Bacillus cereus* using the 16S rRNA method (Fox et al. 1977). The strain was maintained in LB agar at 4°C or in LB containing 20% glycerol at -70°C.

### Culture conditions

The microorganism was cultivated in 500 ml flasks with 100 ml medium containing 1.5% olive oil, 2.5% PEG-400, 0.5% soluble starch, 2% yeast extract  $(NH_4)_2SO_4$ , 0.1%  $K_2PO_4$ , 0.05%  $MgSO_4$ , 0.01%  $CaCl_2$ , and 0.001%  $FeSO_4$ . The initial pH of the culture medium was adjusted to 7.0 with 1 M NaOH. After inoculation with 10 ml of a preculture in the same medium, the culture was grown at 30°C and 220 rpm for 24 h.

### Purification of esterase from *Bacillus cereus*

All procedures were performed at room temperature. Cells from a 24-h culture were harvested by centrifugation (8,000 rpm, 15 min). The cell pellet (10 g) was washed twice with 20 mM pH 7.2 Tris-HCl buffer and disrupted by sonication in same buffer with 1 mM EDTA. Cell debris was eliminated by high speed centrifugation (15,000 rpm, 25 min), one volume ethanol was added to the suspension, which was then left at room temperature for 30 min. The precipitate was eliminated by high speed centrifugation (15,000 rpm, 25 min), then another one volume ethanol was added and the suspension left at room temperature for 30 min. The suspension was centrifuged at 15,000 rpm for 25 min and the pellet was re-dissolved in a minimal volume of 20 mM pH 7.2 Tris-HCl buffer. The enzyme solution was dialyzed against the same buffer for 8 h then centrifuged to eliminate un-dissolved pellet. The enzyme solution was loaded onto a DEAE-Sepharose CL6B column (1.0×15 cm) equilibrated with 20 mM pH 7.2 Tris-HCl.

The flow rate was set at 1.5 ml/min. After washing, a linear 0–1 M NaCl gradient in the same buffer was applied. The active fractions eluted were pooled, dialyzed against 20 mM pH 7.2 Tris-HCl with 1 M  $(NH_4)_2SO_4$ , and applied to a column of Phenyl Sepharose (1.0×5 cm), equilibrated in 20 mM pH 7.2 Tris-HCl containing 1 mM  $(NH_4)_2SO_4$ . After washing, a gradient from 1 M  $(NH_4)_2SO_4$  to 0 M  $(NH_4)_2SO_4$  was applied, the active fraction were pooled, and dialyzed against 10 mM pH 7.2 phosphate buffer. The enzyme solution was concentrated and re-dissolved in 0.5 ml 10 mM pH 7.2 phosphate buffer, and the solution was then loaded onto a Sephacryl S200 HR column (1.0×100 cm) equilibrated and eluted with 10 mM phosphate buffer (pH 7.2). Elution was performed at a rate of 1 ml/min and active fractions were pooled.

### Molecular mass determination by size-exclusion chromatography

A Sephacryl S200 HR column (GE, Piscataway, NJ) was calibrated with the following proteins (Serva, Heidelberg, Germany): ribonuclease (13.7 kDa), ovalbumin (45 kDa), bovine serum albumin (67 kDa), and alcohol dehydrogenase (150 kDa).

### Protein determination

Protein concentration was determined as described (Bradford 1976) with bovine serum albumin as standard protein.

### Gel electrophoresis

SDS-PAGE of the purified enzyme was conducted as described by Laemmli (1970) using 12% gels at pH 8.8. The same procedure was used to set up a calibration curve with marker proteins from the LMW Electrophoresis Calibration Kit supplied by Sigma.

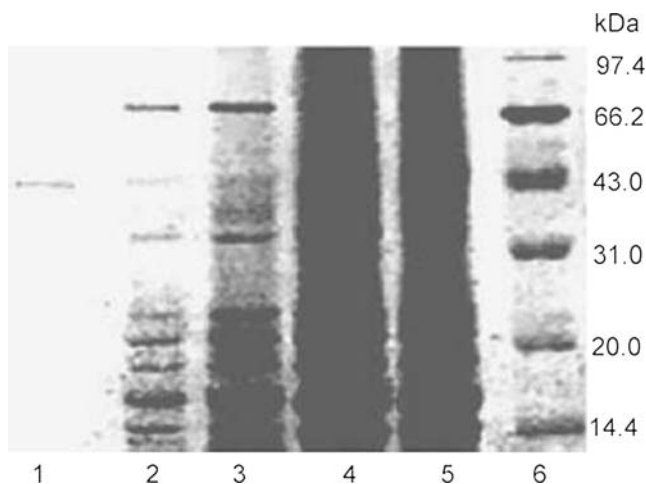
### Determination of N-terminal amino acid sequence

The purified esterase on SDS-PAGE gel was transferred to a PVDF membrane (Bio-Rad, Richmond, CA). After blotting, the PVDF membrane was stained and the esterase band was excised. N-terminal Edman sequencing was performed on an Applied Biosystems Procise 419 sequencer (Foster City, CA).

### Hydrolysis activity assay of racemic

#### 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol

1-Chloro-3-(1-naphthoxy)-2-acetoxypropanol (0.2 mg) was dissolved in ethanol, and 0.85 ml 20 mM phosphate buffer (pH 7.2) was added to a flask with a lid. Enzyme (0.05 mg) was then added and the system was sealed and



**Fig. 2** SDS-PAGE analysis of purified esterase. Lanes: 1 Sephacryl S200 HR, 2 Phenyl sepharose, 3 DEAE-Sepharose CL6B, 4 ethanol precipitation, 5 sonication debris, 6 protein marker

shaken at 30°C, 200 rpm for 6 h. The reaction mixture was extracted with 1 ml ethyl acetate. The ethyl acetate extract (10  $\mu$ l) was applied to a Phenomenex 3022 column (<http://www.phenomenex.com>) and was eluted with a mobile phase of 80% hexane/20% ethyl acetate (v/v). The UV absorption of 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol was monitored at 256 nm. One unit of enzyme activity is defined as the amount of enzyme hydrolyzing 1  $\mu$ mol substrate per minute.

#### Determination of ee value by capillary electrophoresis

Solvent in the sample from the hydrolysis reaction was evaporated, and the residue was submitted to a silicon column mobile phase. The mobile phase was same as for the hydrolysis activity assay. Purified ester was reacted with 20 volumes of 10% NaOH and a two times molar ratio of isopropylamine for 12 h at room temperature. The product was then applied to a Beckman P/ACE SYSTEM-MDQ capillary electrophoresis system (Fullerton, CA) with an unmodified silicon capillary (75  $\mu$ m I.D.  $\times$  375  $\mu$ m O.D.). Capillary electrophoresis analysis was conducted as described (Wang et al. 2001). The enantioselectivity for the remaining substrate was expressed by

enantiomeric excess (ee S) and enantiomeric ratio (*E* value) (Chen et al. 1982):

$$e.e.S = \frac{S_S - S_R}{S_S + S_R} \times 100\%$$

$$E = \frac{\ln[(1-c)(1-eeS)]}{\ln[(1-c)(1+eeS)]}$$

where *c* stands for conversion, and *S<sub>S</sub>* and *S<sub>R</sub>* stand for S-propranolol and R-propranolol.

#### Esterase and activity assay

The standard assay system consisted of 10 mM para-nitrophenyl fatty acid esters, 100 mM NaH<sub>2</sub>PO<sub>4</sub>-citric acid buffer, pH 7.5, 0.1% (w/v) of sodium deoxycholate, and 10  $\mu$ g enzyme, in a final volume of 0.2 ml. The mixture, without the substrate, was brought to 40°C. The reaction was started by the addition of substrate (10  $\mu$ l), which had previously been prepared as a stock solution in isopropanol. Changes in absorbance at 405 nm were recorded for 20 min using a 721 model spectrophotometer (Kebo, Hangzhou, PR China). The reaction was stopped by adding two volumes of ethanol. The enzyme was replaced by water to generate a blank, and, in every measurement, the effect of non-enzymatic hydrolysis of substrates was taken into account and subtracted from the value measured when the enzyme was added. The molar extinction coefficients of para-nitrophenol were also determined under each reaction condition prior to the measurements. One unit of esterase activity is defined as the amount of enzyme forming 1  $\mu$ mol substrate per minute.

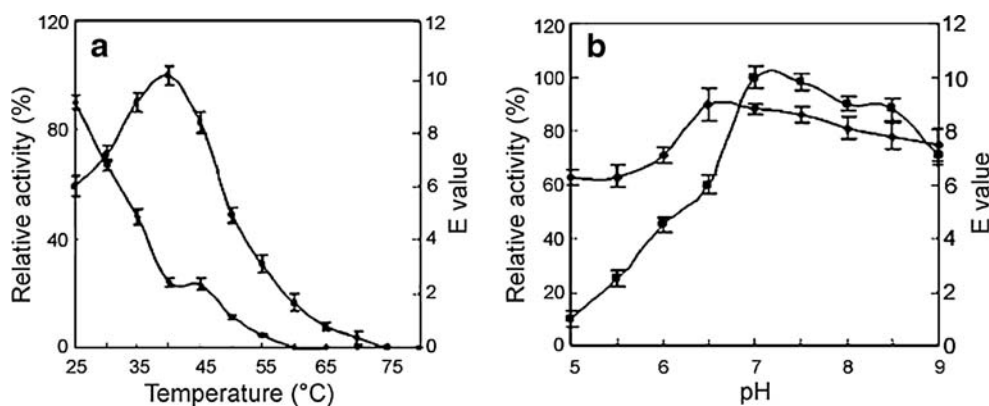
## Results

### Screening

A collection of microorganisms screened from soil samples was used for the resolution of 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol. Whole cells of strain WL 24 were found to have the highest selectivity for resolution of the

**Table 1** Summary of esterase purification

Purification step	Total protein (mg)	Total activity (U)	Specific activity (U/mg protein)	Recovery (%)	Purification fold
Crude extract	786	5.6	0.007	100	1
Ethanol precipitation	116	4.5	0.04	80.0	5.7
DEAE-Sepharose CL6B	9	2.2	0.24	39.2	34.2
Phenyl Sepharose	1.2	0.9	0.75	16	107.1
Sephacryl S200HR	0.1	0.32	3.2	5.7	457.1



**Fig. 3** Temperature (a) and pH (b) optima of purified esterase from *Bacillus cereus*. Left axis (■) relative activities, right axis (◆) E values. **a** Purified esterase (10  $\mu$ g) was assayed under standard hydrolysis assay conditions at various temperatures. **b** Purified esterase (10  $\mu$ g)

was assayed under standard hydrolysis assay conditions with different buffers (100 mM  $\text{NaH}_2\text{PO}_4$ -citric acid buffer for pH 5.0–8.0; 100 mM glycine/NaOH for pH 8.5–9.0)

substrate, while the supernatant of the same culture exhibited no activity towards the substrate; therefore, the enzyme is an intracellular protein. Taxonomic studies identified the strain as *Bacillus cereus*.

#### Esterase purification

The esterase was purified 457 fold with a four-step purification procedure based on the hydrolysis activity assay for 1-chloro-3-(1-naphthoxy)-2-acetoxypropanol. The purified fraction appeared homogenous on SDS gels (Fig. 2, Table 1).

#### Physical properties of the esterase

The molecular mass of the esterase was determined by SDS-PAGE as 43 kDa. The molecular mass of the native enzyme was determined as 42 kDa after calibration of the gel filtration (Sephacryl S200 HR) column. Maximum activity was found at pH 7.0 and 40°C. Maximum enantioselectivity was also found at pH 7.0, with the temperature optimum for enantioselectivity decreasing with increasing transformation temperature (Fig. 3).

**Table 2** Specific activity for hydrolysis of various substrates. *p*-NP Para-nitrophenyl

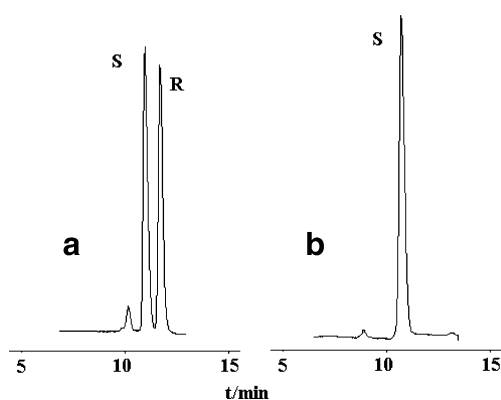
Substrate (carbon chain length)	Specific activity (U/mg)
<i>p</i> -NP acetate (C2)	$116.3 \pm 9.4 \times 10^3$
<i>p</i> -NP butyrate (C4)	$124.6 \pm 7.8 \times 10^3$
<i>p</i> -NP caproate (C6)	$215.6 \pm 8.4 \times 10^3$
<i>p</i> -NP laurate (C12)	$89.5 \pm 2.3 \times 10^3$
<i>p</i> -NP tetradecanoate (C14)	$64.6 \pm 2.1 \times 10^3$
<i>p</i> -NP oleate (C18)	$32.4 \pm 3.5 \times 10^3$
<i>p</i> -NP stearate (C18)	$21.1 \pm 0.8 \times 10^3$

#### Substrate specificity of the esterase

Esterase specificity was tested on various *p*-nitrophenyl esters with increasing chain length. The results reported in Table 2 indicate that esterase activity is influenced strongly by the chain length of the acyl group. Enzyme activity displayed an increasing trend in terms of the rate of ester hydrolysis up to C6, and decreased as the chain length increased further.

#### N-terminal sequence analysis

The N-terminal amino acid sequence of the esterase from *Bacillus cereus* was determined as MILVTQTMKG. This sequence was compared with sequences of *Bacillus* esterase and lipase as deposited in GenBank. However, no signif-



**Fig. 4** a, b S-Propranolol produced by a chemoenzymatic process employing whole cells of *Bacillus cereus* WL 24. **a** Standard propranolol, **b** whole cell resolution: 100 mM, phosphate-trihydroxyethylamine buffer (pH 3.0), 30 mM hydroxypropyl- $\beta$ -cyclodextrin, 50 cm unmodified capillary, 20 kV voltage. The conversion of the substrate was controlled beyond 73%; the ee value reached 99%

icant homology to other lipase or esterase sequences was found.

## Discussion

The use of *Bacillus* carboxylesterase for the production of valuable chiral building blocks seems to be a promising approach. For example, a *Bacillus cereus* strain demonstrated hydrolysis activity towards prochiral 2-phenyl-1,3-propanediol diacetate and accumulated the corresponding chiral monoacetates in the reaction mixture (Mitsui et al. 2007); a novel carboxylesterase with high enantioselectivity toward racemic esters of 1,2-O-isopropylidenglycerol was isolated from *Bacillus coagulans* (Molinari et al. 1996; Romano et al. 2005); an extracellular esterase from *Bacillus sphaericus* hydrolyzed ethyl 2-hydroxyalkanoates with high enantioselectivity (Jackson et al. 1995); resolution of R, S-naproxen enantiomers by hydrolysis of the corresponding racemic ethyl esters was achieved by employing carboxylesterase from *Bacillus* sp. (Quax and Broekhuizen 1994), >while the stereospecific hydrolysis of racemic menthylacetate was achieved using esterase from *Bacillus subtilis* (Brookes and Lilly 1986).

There are many reports describing resolution of the same substrate by lipase, such as lipase from *Pseudomonas* sp., which possesses a high E value (around 100, Bevinakatti and Banerji 1991). The esterase from *Bacillus cereus* had a moderate E value (around 8 at 30°C). However, it was relatively economical when whole cells were used for kinetic resolution of the substrate. If conversion of the substrate was controlled beyond 73%, the ee value could reach 99% (Fig. 4). When 1 g dry cells was used, 73% conversion could be obtained for 0.8 g substrate in 12 h. To our knowledge, this is also the first report of the utilization of *Bacillus cereus* in the resolution of 1-chloro-3-(1-naphthoxy)-2-acetoxypentanol. The lack of homology of the N-terminal sequence to other known esterase sequences indicates that the esterase from *Bacillus cereus* might represent a novel esterase.

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