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Note



Overproduction and Substrate Specificity of 3-Isopropylmalate Dehydrogenase from *Thiobacillus ferrooxidans*

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We constructed an overexpression system in *Escherichia coli* of the *leuB* gene coding for 3-isopropylmalate dehydrogenase in *Thiobacillus ferrooxidans*. *E. coli* harboring the plasmid we constructed, pKK leuB1, produced 17-fold the enzyme protein of the expression system previously used for purification. The substrate specificity of the enzyme was analyzed with synthetic (2R, 3S)-3-alkylmalates. The 3-isopropylmalate dehydrogenase of *Thiobacillus ferrooxidans* had broad specificity toward the alkylmalates.

Key words: Thiobacillus ferrooxidans; 3-isopropylmalate dehydrogenase; leuB; overproduction; substrate specificity

3-Isopropylmalate dehydrogenase (EC 1.1.1.85), a key enzyme in leucine biosynthesis, catalyzes the oxidative decarboxylation of the substrate 3-isopropylmalate to 2-oxoisocaproate simultaneously with dehydrogenation. The enzyme has been found in a wide variety of bacteria and the amino acid sequences of the enzyme from various microorganisms including yeasts are similar. ¹⁻¹⁰⁾ The catalytic mechanism of 3-isopropylmalate dehydrogenase may resemble that of bacterial isocitrate dehydrogenase. The X-ray crystal structure of *Thermus thermophilus* 3-isopropylmalate dehydrogenase. ¹¹⁻¹³⁾ is like that of *Escherichia coli* isocitrate dehydrogenase. ^{14,15)} Most of the residues that seem from crystallographic results to be key to enzyme function are found in both enzymes.

We previously cloned the *leuB* gene coding for the 3-isopropylmalate dehydrogenase of an acidophilic chemolithotrophic bacterium, *Thiobacillus ferrooxidans*, in *E. coli* and purified the enzyme to homogeneity from *E. coli* cells harboring a recombinant plasmid, pTFL101, containing the *leuB* gene. The properties of the enzyme are similar to those of the *Salmonella typhimurium* enzyme, secept for substrate specificity. The *T. ferrooxidans* 3-isopropylmalate dehydrogenase utilizes malate as a substrate as well as 3-isopropylmalate. The enzymes from an extreme thermophile, *T. thermophilus*, and thermoacidophilic archaeon, *Sulfolobus* sp. strain 7, also utilize the various 3-alkylmalates with broad substrate specificity.

The purification of the *T. ferrooxidans* 3-isopropylmalate dehydrogenase from the *E. coli* transformant was

not enough for further investigation, for example, crystallography, because the *E. coli* was grown in minimal medium at a low cell yield. Because of the high demand for 3-isopropylmalate dehydrogenase, we undertook to construct an overexpression system. In this paper, we describe an construction of the overexpression plasmid and report the substrate specificity of the 3-isopropylmalate dehydrogenase from *T. ferrooxidans* purified from *E. coli* using the overexpression system.

The T. ferrooxidans leuB structural gene was amplified by PCR with the synthetic oligonucleotides 5'-GGAGAATTCATGAAAAAAATGGCC-3' and 5'-ATCCCCGGGTCAATCCTTCAGGTT-3'. The underlined sequences are restriction sites of EcoRI and SmaI, respectively. The amplified fragment was blunted with DNA polymerase I and phosphorylated with ATP and polynucleotide kinase. The phosphorylated fragment was ligated with pKK223-3 (Pharmacia LKB Biotechnology Inc.), which had been digested with SmaI, and then dephosphorylated. The ligated plasmid was digested with Eco RI and ligated with itself, giving pKK leuB1. E. coli JA221 harboring pKKleuB1 had high 3isopropylmalate dehydrogenase activity: 17-fold that of E. coli HB101 harboring pTFL101 (Table I). Earlier, we purified the enzyme by three kinds of column chromatography, DEAE-Toyopearl 650 M, butyl-Toyopearl 650 M (Tosoh Corp.), and hydroxyapatite. 17) However, with this overexpression system, purification was done with only two kinds of column chromatography, DEAE-Toyopearl 650 M and Q-Sepharose Fast Flow (Pharmacia LKB Biotechnology Inc.) (data not shown).

The T. ferrooxidans 3-isopropylmalate dehydrogenase utilizes malate as a substrate as well as 3-isopropylmalate. The substrate specificity of the enzyme was analyzed with (2R, 3S)-3-alkylmalates synthesized as described previously. Table II lists the kinetic constants K_m and k_{cat} for several alkylmalates of T. ferrooxidans 3-isopropylmalate dehydrogenase. All of these alkylmalates served as a substrate of the enzyme, so its substrate specificity seemed to be broad. The K_m value decreased with increasing hydrophobicity of the alkyl group at position 3 up to 3-ethylmalate and increased afterwards, and k_{cat}/K_m increased up to 3-ethylmalate and decreased afterwards. This result coincides with those for enzymes from T. thermophilus on T and T and T and T and T are relationship between the cata-

Strain (plasmid)	Medium	IPTG ^a (1 mm)		Specific activity ^b (units/mg)		Total activity ^c (units)	
JA221 (pKKleuB1)	LB	+		8.56		810	
JA221 (pKKleuB1)	$\mathbf{L}\mathbf{B}$			5.65		514	
JA221 (pKKleuB1)	minimal	-		6.74		150	
JA221 (pTFL101)	minimal			2.53		49.6	
JA221 (pTFL101)	LB	+		0.12		7.71	
HR101 (nTFI 101)	minimal	_		2 84		48 4	

Table I. 3-Isopropylmalate Dehydrogenase Activity of E. coli Transformants

Table II. Kinetic Parameters for the Catalysis of (2R, 3S)-3-Alkylmalate by *Thiobacillus ferrooxidans* 3-Isopropylmalate Dehydrogenase

π^{a}	K _m (mm)	k_{cat} (s ⁻¹)	$\frac{k_{\rm cat}/K_{\rm m}}{({\rm s}^{-1}\cdot{\rm mM}^{-1})}$
0	7.7	5.0	0.65
0.50	0.050	28	560
1.00	0.013	38	2900
1.30	0.015	28	1900
1.98	0.074	2.7	36
2.30	0.11	20	180
	0 0.50 1.00 1.30 1.98	π (mM) 0 7.7 0.50 0.050 1.00 0.013 1.30 0.015 1.98 0.074	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$

^a The π values are the Hansch constants for the alkyl substituents.²²⁾

^b Malate has only one asymmetric carbon atom.

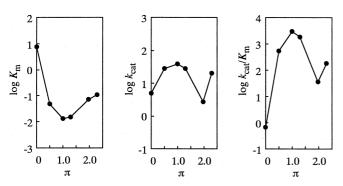


Fig. 1. Relationship between the Hydrophobicity of the Alkyl Group and $K_{\rm m}$, $k_{\rm cat}$, and $k_{\rm cat}/K_{\rm m}$ for the Catalysis of Alkylmalate by 3-Isopropylmalate Dehydrogenase.

The data were replotted from the values in Table II. Energies are in kcal/mol. The π values are the Hansch constants for the alkyl substituents.²²⁾

lytic properties and the hydrophobicity of the alkyl group is shown in Fig. 1. These results suggest that the reaction was generally independent of the hydrophobicity. The *T. ferrooxidans* 3-isopropylmalate dehydrogenase had broad substrate specificity toward alkylmalates, but it had no activity with isocitrate, ¹⁷⁾ which has a negatively charged carboxymethyl group instead of an alkyl group. This finding suggests a chemical difference in the recognition sites of the 3-isopropylmalate de-

hydrogenase and the isocitrate dehydrogenase for the substituent at C-3 of malate.

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a Isopropyl-β-D-thiogalactopyranoside.

^b Specific activity is defined as μ mol of NADH formed per milligram of protein per minute.

^c Total activity is defined as activity per 100 ml of culture.