

Purification and some properties of (1*R*,2*S*)-1,2-dihydroxy-3,5-cyclohexadiene-1,4-dicarboxylate dehydrogenase from *Comamonas testosteroni* T-2

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Abstract

Inducible (1*R*,2*S*)-1,2-dihydroxy-3,5-cyclohexadiene-1,4-dicarboxylate (diene-diol) dehydrogenase was found in extracts of *Comamonas testosteroni* T-2 grown in *p*-toluate-or terephthalate-salts medium and it was purified using anion exchange, hydrophobic interaction and gel filtration chromatography. The enzyme is a homodimer with subunit M_r 39 000. It had a specific activity of 500 mkat/kg of protein and was activated by the addition of Fe^{2+} . The dehydrogenase converted 1 mol diene-diol and 1 mol NAD^+ to 1 mol protocatechuic acid, 1 mol NADH and 1 mol CO_2 . Apparent K_m -values of 43 μM (NAD^+) and about 90 μM (diene-diol) were determined. The hydride ion was transferred to the *si* face of NAD^+ .

Keywords: Terephthalate; Diene-diol dehydrogenase; Enzyme purification; *Comamonas testosteroni* T-2; Biodegradation

1. Introduction

Phthalate, *iso*-phthalate and terephthalate (1,2-, 1,3- and 1,4-dicarboxybenzene, respectively) are unreactive aromatic compounds that can enter the environment from industrial or natural sources [1,2]. Bacterial degradation of all three isomers has been reported [1,3,4] but the metabolism of phthalate itself is best understood [5]. In contrast to the three steps from phthalate to protocatechuic acid, there are only two steps from terephthalate to this intermediate (Fig. 1).

Whereas the multi-component terephthalate dioxygenase (Fig. 1) has been partially characterized [2], the (1*R*,2*S*)-1,2-dihydroxy-3,5-cyclohexadiene-1,4-dicarboxylate (diene-diol) dehydrogenase has not been described. We now report the purification and some properties of this dehydrogenase, the last step specific to the degradation of *p*-toluate to protocatechuic acid in *Comamonas testosteroni* T-2 [6].

2. Materials and methods

2.1. Materials

The diammonium salt of the diene-diol was kindly provided by Celgene (Warren, NJ). We identified the

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diene-diol indirectly, using the characteristic of these compounds that water is lost on acidification [7]; the product was derivatized with diazomethane and identified by GC-MS [2,8] as a dimethylated derivative of 2-hydroxyterephthalate. The preparation of the diene-diol was initially chromatographically pure but it deteriorated over time. Complete enzymic transformation of this diene-diol with excess of NAD^+ yielded 0.5 mol protocatechuate as the sole product; the impurity was not altered. We concluded that the preparation used contained 50% (wt/wt) diene-diol. $[4\text{-}^3\text{H}]\text{-NAD}^+$ was prepared from $\text{D-}[1\text{-}^3\text{H}]\text{-glucose}$ [9,10]. The Carbosieve SII column (100/120 mesh; $3\text{ m} \times 2\text{ mm}$; stainless steel) was from Supelco (Bellefonte, PA). All other chemicals and materials used were from sources indicated elsewhere [2,8,11].

2.2. Organism, growth conditions and preparation of cell-free extracts

Comamonas testosteroni T-2 (DSM 6577 [12]) was grown in toluate- or terephthalate-salts medium as described previously [2,8]. Cells harvested in the mid-log growth phase were disrupted in a French pressure cell to obtain crude extract [2].

Diene-diol dehydrogenase activity was measured photometrically at 340 nm as diene-diol-dependent generation of NADH in an assay containing (in 1 ml) 40 μmol Tris sulfate, pH 7.5, 600 nmol NAD^+ and 10 to 100 μg of protein. The reaction was started by the addition of 750 nmol diene-diol [2]. If required, the chelator EDTA (up to 125 μM) was added to the

reaction, sometimes with preincubation. The standard reaction, in 8-ml septum vials, was used to follow release of CO_2 from diene-diol; samples (0.3 ml) of the gas phase were taken by gas-tight syringe and examined by GC with thermal conductivity detection. The organic product from diene-diol was tentatively identified and quantified by reversed-phase HPLC at room temperature [2,13]. The M_r of the native enzyme was estimated by gel filtration through a Superose 6 column [14]. SDS-PAGE was used for the determination of M_r under denaturing conditions and to monitor the protein purification. Proteins in gels were visualized by staining with Coomassie brilliant blue (the method of Laemmli) or silver [15]. Protein concentration was measured by the method of Bradford with bovine serum albumin as standard. The stereospecificity of the diene-diol dehydrogenase was determined in the standard assay (see above and [2]), containing 0.4 nmol $[4\text{-}^3\text{H}]\text{NAD}^+$ (170 000 dpm). After completion of the reaction, 10 U of yeast alcohol dehydrogenase (Sigma) and 2 μl acetaldehyde were added to the reaction to reoxidize the generated NADH in a defined stereospecific (pro-*R*) reaction [9]. The reaction products were examined by reversed-phase HPLC and in a liquid scintillation counter as described elsewhere [9].

2.3. Purification of the diene-diol dehydrogenase

The diene-diol dehydrogenase was routinely purified from toluate-grown cells because the enzyme

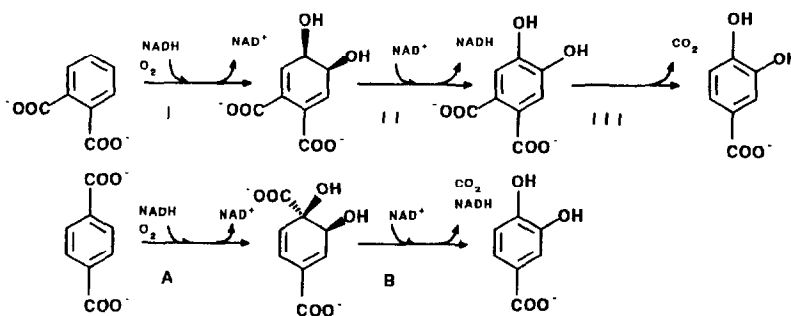


Fig. 1. Initial steps in the degradative pathways of phthalate and terephthalate. The phthalate dioxygenase system (reaction I) is one of the best-characterized dioxygenase systems [26]. The oxidation of the diene-diol to 4,5-dihydroxyphthalate (reaction II) has not been studied in detail, whereas the enzyme catalyzing decarboxylation to protocatechuate (reaction III) has been purified from two organisms [27,28]. The terephthalate dioxygenase system (reaction A) has been partially characterized [2] and the diene-diol dehydrogenase (reaction B) was detected in our preliminary experiments (cited in [6]): the reactions have also been detected in *Pseudomonas putida* (cited in [6]).

cochromatographed on two columns (see below) with component A (oxygenase) of 4-sulfobenzoate 1,2-dioxygenase system [16], which is expressed in terephthalate-but not in toluate-grown cells (Schläfli Oppenberg, 1995, unpublished data). 20 mM Tris sulfate, pH 7.5, containing 1 mM dithiothreitol (buffer A), was used as basis for all eluents. The purification procedure is, in most practical aspects, similar to that of the terephthalate dioxygenase system [2].

Step 1: the preparation of crude extracts free of nucleic acids [2,16], which were removed by protamine sulfate precipitation.

Step 2: anion-exchange chromatography on a Mono Q HR 10/10 (Pharmacia, 10 × 100 mm) with a gradient of Na₂SO₄ as described for the terephthalate dioxygenase system [2]. The diene-diol dehydrogenase activity eluted at the beginning of the gradient, after fraction R (containing the reductase function of the terephthalate dioxygenase system) and overlapping with any component A of the 4-sulfobenzoate dioxygenase system. Fractions containing significant diene-diol dehydrogenase activity were pooled and concentrated by ultrafiltration (Centriprep, 10 kDa exclusion limit).

Step 3: hydrophobic interaction chromatography was done with a Phenyl-Superose HR 5/5 column (5 × 50 mm, Pharmacia) [cf. 2]. The diene-diol dehydrogenase from step 2 was loaded on to the column and a gradient of (NH₄)₂SO₄ in buffer A was applied which decreased over the course of 30 min from 0.5 M to 0 M. A single diene-diol dehydrogenase activity eluted at the end of the gradient which was essentially pure when the purification was accomplished from toluate-extract. The diene-diol dehydrogenase from terephthalate-extract was contaminated after this step with oxygenase A from the 4-sulfobenzoate dioxygenase system. Crude diene-

diol dehydrogenase was pooled and concentrated (Centricon, see above).

Step 4: a Superose 6 column (10 × 300 mm, Pharmacia) was equilibrated (flow of 0.5 ml min⁻¹) on buffer A, containing 0.2 M Na₂SO₄. The diene-diol dehydrogenase (200 μl) from step 3 was loaded onto the column and diene-diol dehydrogenase activity eluted after 30 min in a symmetrical peak, containing essentially one protein as monitored by SDS-PAGE.

3. Results and discussion

The diene-diol dehydrogenase was detected in extracts of cells of *C. testosteroni* T-2 which could utilize terephthalate (e.g. toluate-or terephthalate-grown) and was not observed under other conditions (e.g. succinate-grown cells). We consider the dehydrogenase to be an inducible enzyme. We obtained 90 μg of pure enzyme from about 54 mg protein in the extract (Table 1). Given a purification factor of 55, the amount of diene-diol dehydrogenase in the cell is calculated to be about 2% of total soluble protein. The amount of the terephthalate dioxygenase system is in the same range [cf. 2]. The diene-diol dehydrogenase activity observed in the extract (10 mkat/kg of protein; Table 1) is five times the activity needed to support the growth rate observed in terephthalate-salts medium [2].

The diene-diol dehydrogenase could be purified to homogeneity in three steps (Fig. 2, Table 1) and an *M_r*-value of 39 000 was observed under denaturing conditions (Fig. 2). The *M_r* obtained under native conditions was 60 000 ± 10 000. These values suggest a homodimeric structure. The known diol dehydrogenases are described as dimeric [14], tetrameric

Table 1
Purification of the diene-diol dehydrogenase from *C. testosteroni* T-2

Protein fraction	Volume (ml)	Total protein (μg)	Total activity (nkat)	Specific activity (mkat/kg of protein)	Yield (%)	Purification (fold)
Crude extract	0.91	53 700	518	9.6	100	1
Anion exchange chromatography	3.63	4200	109	25.8	21	2.7
Hydrophobic interaction chromatography	0.93	128	51	400	9.8	41.5
Gel filtration chromatography	2.0	90	48	533	9.3	55.3

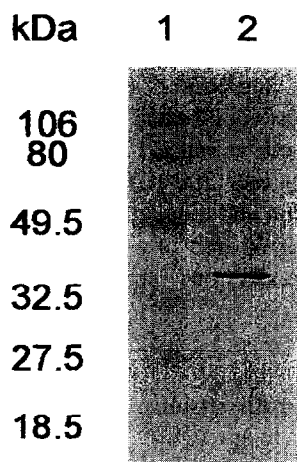


Fig. 2. SDS-PAGE of purified diene-diol dehydrogenase. Proteins in the gel were visualized by silver-staining. 1, prestained low-range marker proteins (Bio-Rad); 2, 1 μg of the diene-diol dehydrogenase purified by gel filtration chromatography.

[17–22] or hexameric [23], with M_s s of about 25 000 to 30 000 [14,18–20,22,23], 50 000 [21] or 110 000 [17]. The diol dehydrogenase from *C. testosteroni* T-2 therefore appears to be unusual both in subunit- M_r and subunit composition.

Enzyme catalysis gave a 98% yield of NADH from NAD^+ and a 93% yield of protocatechuate from NAD^+ . The anticipated gaseous product of the reaction, CO_2 , was detected in the gas phase in 57–83% of the theoretical yield by GC. This product, which was absent from control experiments, had the same retention time as authentic material. The data indicate that 1 mol diene-diol and 1 mol NAD^+ yield 1 mol protocatechuate, 1 mol NADH, and 1 mol CO_2 .

The enzyme was specific for NAD^+ , NADP^+ was not a cofactor. The apparent K_m for NAD^+ (43 μM) was low, similar to that for benzene dihydrodiol dehydrogenase [17], whereas other diene-diol dehydrogenases have values ranging up to about 900 μM . The apparent K_m for the diene-diol, about 90 μM , is at the higher end of the range observed, values $< 2 \mu\text{M}$ being known [18]. Addition of FAD or FMN had no effect on the enzyme activity.

We examined the stereospecificity of hydride transfer by the diene-diol dehydrogenase to $[4\text{-}^3\text{H}]\text{NAD}^+$. In experiments with different enzyme preparations we found that the secondary enzyme,

yeast alcohol dehydrogenase, which is specific for the pro-*R* hydride of NADH, removed more than 85% of the tritium label from the $[4\text{-}^3\text{H}]\text{NADH}$ generated by the diene-diol dehydrogenase. We therefore conclude that the diene-diol dehydrogenase specifically transfers hydride ions to the (4*S*)-position (*si*-face) of NAD^+ .

We observed the highest activity of the diene-diol dehydrogenase at pH 8.2 in 20 mM Tris-sulfate. Two described diol dehydrogenases have a similar optimal pH [17,22], and most other enzymes a significantly higher optimal pH (pH 9.0 and more) [18–21,23].

The UV-visible spectrum of the enzyme, with no absorbance in the visible or near UV (a λ_{max} at 280 nm) gave no evidence for prosthetic groups with chromophores. The reaction was activated by a factor of about two on the addition of iron (saturation at 60 μM Fe^{2+}). The reaction could be totally inhibited by the addition of EDTA, the amount of which decreased with increasing preincubation. Addition of iron could overcome the inhibition.

In a recent review, Reid and Fewson [24] classified the microbial alcohol dehydrogenases into three major categories, the first category comprising NAD(P)-dependent enzymes in three sub-groups. The known properties of the diene-diol dehydrogenase do not fit the characteristics of any of these sub-groups. Group I enzymes are pro-*R* specific, zinc-dependent enzymes consisting of about 350 amino acid residues. Some aromatic diene-diol dehydrogenases can be attributed to group II, which consists mainly of smaller (25–30 kDa), metal-independent homomultimeric enzymes with predominant pro-*S* specificity and distinct sequence similarities [24,25]. The diene-diol dehydrogenase would be an unusual member of group II for its subunit size and iron requirement. However, determination of an apparent M_r by SDS-PAGE does not yield absolute values and the native M_r could indicate a smaller subunit size. Group III of NAD(P)H requiring alcohol dehydrogenases contains long-chain enzymes (40–100 kDa), that can be activated by iron. No aromatic diol dehydrogenase has been attributed to this group of alcohol dehydrogenases so far, but the diene-diol dehydrogenase and also *cis*-benzene dihydrodiol dehydrogenase could be members of this group. With respect to the known enzymes of this reaction type,

the diene-diol dehydrogenase shows most similarity to benzene dihydrodiol dehydrogenase (pH optimum, K_m (NAD⁺), activation by Fe²⁺) [17].

Acknowledgements

H.R.S.O. was supported by a grant from the Swiss Federal Institute of Technology, Zürich; H.L. was supported by funds from the University of Konstanz.

References

- [1] Keyser, P., Pujar, B.G., Eaton, R.W. and Ribbons, D.W. (1976) Biodegradation of phthalates and their esters by bacteria. *Environ. Health Perspect.* 18, 159–166.
- [2] Schläfli, H.R., Weiss, M., Leisinger, T. and Cook, A.M. (1994) Terephthalate 1,2-dioxygenase system from *Comamonas testosteroni* T-2: purification and some properties of the oxygenase component. *J. Bacteriol.* 176, 6644–6652.
- [3] Ribbons, D.W., Keyser, P., Kunz, D.A., Taylor, B.F., Eaton, R.W. and Anderson, B.N. (1984) Microbial degradation of phthalates. In: *Microbial Degradation of Organic Compounds* (Gibson, D.T., Ed.), pp. 371–397. Dekker, New York.
- [4] Ribbons, D.W., (1988) More about phthalates. In: *Microbial Catabolism and the Carbon Cycle* (Hagedorn, S.R., Hanson, R.S. and Kunz, D.A., Eds.), pp. 85–100. Harwood, Chur.
- [5] Batic, C.J., LaHaie, E. and Ballou, D.P. (1987) Purification and characterization of phthalate oxygenase and phthalate oxygenase reductase from *Pseudomonas cepacia*. *J. Biol. Chem.* 262, 1510–1518.
- [6] Locher, H.H., Malli, C., Hooper, S.W., Vorherr, T., Leisinger, T. and Cook, A.M. (1991) Degradation of *p*-toluic acid (*p*-toluenecarboxylic acid) and *p*-toluenesulphonic acid via oxygenation of the methyl sidechain is initiated by the same set of enzymes in *Comamonas testosteroni* T-2. *J. Gen. Microbiol.* 137, 2201–2208.
- [7] Whited, G.M., McCombie, W.R., Kwart, L.D. and Gibson, D.T. (1986) Identification of *cis*-diols as intermediates in the oxidation of aromatic acids by a strain of *Pseudomonas putida* that contains a TOL plasmid. *J. Bacteriol.* 166, 1028–1039.
- [8] Locher, H.H., Leisinger, T. and Cook, A.M. (1991) 4-Toluene sulfonate methyl-monooxygenase from *Comamonas testosteroni* T-2: purification and some properties of the oxygenase component. *J. Bacteriol.* 173, 3741–3748.
- [9] Schläfli, H.R., Baker, D.P., Leisinger, T. and Cook, A.M. (1995) Stereospecificity of hydride removal from NADH by reductases of multi-component non-heme iron oxygenase systems. *J. Bacteriol.* 177, 831–834.
- [10] Wermuth, B., Münch, J.D.B. and von Wartburg, J.P. (1979) Stereospecificity of hydrogen transfer of aldehyde reductase. *Experientia* 1288–1289.
- [11] Bünz, P.V. and Cook, A.M. (1993) Dibenzofuran 4,4a-dioxygenase from *Sphingomonas* sp. strain RW1: angular dioxygenation by a 3-component enzyme system. *J. Bacteriol.* 175, 6467–6475.
- [12] Thurnheer, T., Köhler, T., Cook, A.M. and Leisinger, T. (1986) Ortho-nitrobenzoic acid and analogues as carbon sources for bacteria: growth physiology and enzymic desulfonation. *J. Gen. Microbiol.* 132, 1215–1220.
- [13] Locher, H.H., Leisinger, T. and Cook, A.M. (1989) Degradation of *p*-toluenesulphonic acid via sidechain oxidation, desulphonation and *meta* ring cleavage in *Pseudomonas (Comamonas) testosteroni* T-2. *J. Gen. Microbiol.* 135, 1969–1978.
- [14] Neidle, E., Shapiro, M.K. and Ornston, L.N. (1987) Cloning and expression in *Escherichia coli* of *Acinetobacter calcoaceticus* genes for benzoate degradation. *J. Bacteriol.* 169, 5496–5503.
- [15] Bloom, H., Beyer, H. and Gross, H.S. (1987) Improved silver staining of plant proteins, RNA and DNA in polyacrylamide gels. *Electrophoresis* 8, 93–99.
- [16] Locher, H.H., Leisinger, T. and Cook, A.M. (1991) 4-Sulphobenzoate 3,4-dioxygenase: purification and properties of a desulphonative two-component enzyme system from *Comamonas testosteroni* T-2. *Biochem. J.* 274, 833–842.
- [17] Axcell, B.C. and Geary, P.J. (1973) Purification and some properties of the enzyme *cis*-1,2-dihydroxy-cyclohexa-3,5-diene (nicotinamide adenine dinucleotide) oxidoreductase (*cis*-benzene glycoldehydrogenase). *Biochem. J.* 136, 927–934.
- [18] Rogers, J.E. and Gibson, D.T. (1977) Purification and properties of *cis*-toluene dihydrodiol dehydrogenase from *Pseudomonas putida*. *J. Bacteriol.* 130, 1112–1124.
- [19] Patel, T.R. and Gibson, D.T. (1974) Purification and properties of (+)-*cis*-naphthalene dihydrodiol dehydrogenase of *Pseudomonas putida*. *J. Bacteriol.* 119, 879–888.
- [20] Nagao, K., Takizawa, N. and Kiyohara, H. (1988) Purification and properties of *cis*-phenanthrene dihydrodiol dehydrogenase in *Alcaligenes faecalis* AFK2. *Agric. Biol. Chem.* 52, 2621–2623.
- [21] Eberspächer, J. and Lingens, F. (1978) Reinigung und Eigenschaften von zwei Chloridazondihydrodiol-Dehydrogenasen aus Chloridazon-abbauenden Bakterien. *Hoppe-Seyler's Z. Physiol. Chem.* 359, 1323–1334.
- [22] Reiner, A.M. (1972) Metabolism of aromatic compounds in bacteria: purification and properties of the catechol-forming enzyme, 3,5-cyclohexadiene-1,2-diol-1-carboxylic acid (NAD⁺) oxidoreductase (decarboxylating). *J. Biol. Chem.* 247, 4960–4965.
- [23] Simpson, H.D., Green, J. and Dalton, H. (1987) Purification and some properties of a novel heat-stable *cis*-toluene dihydrodiol dehydrogenase. *Biochem. J.* 244, 585–590.
- [24] Reid, M.F. and Fewson, C.A. (1994) Molecular characterization of microbial alcohol dehydrogenases. *Crit. Rev. Microbiol.* 20, 13–56.
- [25] Neidle, E., Hartnett, C., Ornston, L.N., Bairoch, A., Rekik, M. and Harayama, S. (1992) *Cis*-diol dehydrogenases encoded by the TOL pWWO plasmid *xy/L* gene and the

- Acinetobacter calcoaceticus* chromosomal *benD* gene are members of the short-chain alcohol dehydrogenase superfamily. *Eur. J. Biochem.* 204, 113–120.
- [26] Batic, C.J., Ballou, D.P. and Correll, C.C., (1992) Phthalate dioxygenase reductase and related flavin-iron-sulfur containing electron transferases. In: *Chemistry and biochemistry of flavoenzymes* (Müller, F., Ed.), pp. 543–556, CRC Press, Boca Raton.
- [27] Pujar, B.G. and Ribbons, D.W. (1985) Phthalate metabolism in *Pseudomonas fluorescens* PHK: purification and properties of 4,5-dihydroxyphthalate decarboxylase. *Appl. Environ. Microbiol.* 49, 374–376.
- [28] Nakazawa, T. and Hayashi, E. (1978) Phthalate and 4-hydroxyphthalate metabolism in *Pseudomonas testosteroni*: purification and properties of 4,5-dihydroxyphthalate decarboxylase. *Appl. Environ. Microbiol.* 36, 264–269.