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## Substrate Specificities of the Soluble and Particulate Methane Mono-oxygenases of *Methylosinus trichosporium* OB3b

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The intracellular location of methane mono-oxygenase (MMO) (soluble or particulate) in *Methylosinus trichosporium* OB3b is dependent on the availability of copper in the growth medium. Raising the  $\text{Cu}^{2+}$  concentration from 1  $\mu\text{M}$  to 5  $\mu\text{M}$  effected a transition from soluble to particulate MMO activity, and changes in major cell polypeptides were observed on SDS-polyacrylamide gels. Organisms containing soluble MMO oxidized a wide range of substrates including *n*-alkanes, *n*-alkenes, aromatic and alicyclic compounds. By contrast, organisms containing particulate MMO did not oxidize aromatic or alicyclic compounds. These observations provide further evidence that the two types of MMO are fundamentally different.

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### INTRODUCTION

The oxidation of methane to methanol by methanotrophs is catalysed by methane mono-oxygenase (MMO). Scott *et al.* (1981*a, b*) discovered that *Methylosinus trichosporium* OB3b possesses soluble or particulate (membrane-bound) cell-free MMO activity depending on the conditions under which it is grown. Particulate MMO was present in organisms grown in continuous culture under oxygen-limited, nitrate-excess conditions, whilst soluble MMO was present in nitrate-depleted cultures. It was suggested therefore, that the availability of nitrate might determine the intracellular location of MMO (Scott *et al.*, 1981*a, b*). Stanley *et al.* (1983) have now shown that in *Methylococcus capsulatus* (Bath) and *M. trichosporium* OB3b the intracellular location of MMO is determined by the availability of copper; particulate cell-free MMO activity was observed when the organisms were grown at high  $\text{Cu}^{2+}$ :biomass ratios, whereas organisms subjected to copper deficiency contained soluble MMO. The growth conditions used by Scott *et al.* (1981*a*) masked the underlying copper effect. The particulate and soluble MMOs were found to differ in inhibitor specificity, the former being more sensitive to metal-chelating agents, although NAD(P)H was found to be the only suitable electron donor for both activities (Scott *et al.*, 1981*a*).

The ability of MMOs from a number of methanotrophs to oxidize a wide range of substrates is well documented (Colby *et al.*, 1977; Stirling *et al.*, 1979; Higgins *et al.*, 1979; Patel *et al.*, 1980). These substrates include *n*-alkanes, *n*-alkenes and ethers, and aromatic, alicyclic and heterocyclic compounds. *Methylomonas methanica*, however, appears to have a more restricted substrate range. The particulate cell-free MMO from this organism does not oxidize aromatic, alicyclic and heterocyclic compounds (Stirling *et al.*, 1979).

This work compares directly the substrate specificities of *M. trichosporium* OB3b cells containing either soluble or particulate MMO.

### METHODS

*Organism and growth conditions.* *Methylosinus trichosporium* OB3b was obtained from Professor R. Whittenbury, Department of Biological Sciences, University of Warwick, UK. The organism was grown on the nitrate mineral salts medium (NSM) of Cornish *et al.* (1984). This contains 1  $\mu\text{M}$ - $\text{Cu}^{2+}$  and is here referred to as the 'low copper medium'. The 'high copper medium' referred to in the text contains 5  $\mu\text{M}$ - $\text{Cu}^{2+}$ . *M. trichosporium* OB3b was grown

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*Abbreviation:* MMO, methane mono-oxygenase.

on the low copper medium in continuous culture (oxygen-limited) in a 10 l capacity LKB 1601 Ultroform fermenter with methane as sole carbon source (temperature, 30 °C; impeller speed, 500 r.p.m.; pH 6.8; methane flow rate, 250 ml min<sup>-1</sup>; air flow rate, 1000 ml min<sup>-1</sup>; dilution rate, 0.04 h<sup>-1</sup>; biomass density, 1.6–1.8 mg dry wt ml<sup>-1</sup>). The organism was grown on the high copper medium in continuous culture (oxygen-limited) in a 2 l capacity fermenter (L. H. Engineering Co., type L.H.E. CC1500) with methane as sole carbon source (temperature, 30 °C; impeller speed, 400 r.p.m.; pH 6.8; methane flow rate, 50 ml min<sup>-1</sup>; air flow rate, 250 ml min<sup>-1</sup>; dilution rate, 0.04 h<sup>-1</sup>; biomass density, 1.6–1.8 mg dry wt ml<sup>-1</sup>).

**Preparation of suspensions of *M. trichosporium* OB3b.** Organisms were harvested by centrifugation (4420 g, 20 min, 4 °C), washed once with 20 mM-sodium phosphate buffer containing 5 mM-MgCl<sub>2</sub> (pH 7.0) and resuspended in the same buffer to give the required bacterial density (see below). Suspensions prepared in this way were used throughout this work.

**Assay of cell-free MMO activity.** Cell-free extracts were prepared from suspensions of *M. trichosporium* OB3b (30–35 mg dry wt ml<sup>-1</sup>) using sonication to disrupt the cells (4 × 15 s bursts, 16 µm amplitude, with 1 min cooling on ice between bursts). Particulate and soluble fractions were prepared and stored as described by Scott *et al.* (1981a). MMO activity was measured by following the epoxidation of propene, a substrate of soluble and particulate MMO (Colby *et al.*, 1977; Stirling *et al.*, 1979; Scott *et al.*, 1981a). Assays were carried out in 10 ml conical flasks (capped with Suba-seals) which contained 1 ml soluble or particulate fractions (5–10 mg protein). Propene gas (2 ml) was introduced by injection and the reaction was started by adding NADH (5 mM) as electron donor. Flasks were incubated at 30 °C for 5 min in a reciprocating water bath shaker operating at 160 oscillations min<sup>-1</sup>. Propene oxide was measured by gas chromatography using a Porapak Q column (2.1 m × 4 mm i.d.) attached to a Pye Unicam series 204 or 304 instrument fitted with flame ionization detectors [oven temperature, 140 °C; injector, 200 °C; detector, 200 °C; carrier gas (N<sub>2</sub>) flow rate, 40 ml min<sup>-1</sup>].

**Assay of whole-cell oxidations.** Incubations were carried out in 10 ml conical flasks (capped with Suba-seals) containing 1 ml cell suspension (10–12 mg dry wt). Assays were started by injecting the substrate (2 ml gaseous substrate, 2 µl liquid substrate). Sodium formate (20 mM) was used as a source of reducing equivalents in all assays. Flasks were shaken at 30 °C in a reciprocating water bath operating at 160 oscillations min<sup>-1</sup>. Samples were withdrawn for product determination 4 min after starting the incubation and thereafter at 10–12 min intervals over a period of 60 min. Products were measured by gas chromatography using a Pye-Unicam series 204 or 304 instrument fitted with flame ionization detectors [detector, 200 °C; injector, 200 °C; carrier gas (N<sub>2</sub>) flow rate, 40 ml min<sup>-1</sup>]. Products of *n*-alkane oxidation (primary and secondary alcohols, and aldehydes) were analysed using a Porapak Q column (2.1 m × 4 mm i.d.). The column temperatures used for analysis of the oxidation products of different *n*-alkanes were as follows: ethane, 140 °C; propane, 180 °C; butane, 200 °C; pentane, 220 °C and hexane, 220 °C. Products resulting from the oxidation of aromatic or alicyclic compounds were extracted into ethyl acetate (1:1, v/v) and were then analysed using an SP2100 column (2.1 m × 4 mm i.d.). The column temperatures used to analyse oxidation products of aromatic and alicyclic substrates were as follows: benzene, 110 °C ethylbenzene, 100 °C; styrene, 120 °C and cyclohexane, 100 °C.

Ethanol disappearance was measured using gas chromatography. Reactions were carried out in 10 ml conical flasks containing: cell suspension (2.5–10 mg dry wt ml<sup>-1</sup>), 1 ml; and ethanol (15 mM). The flasks were capped with Suba-seals and shaken (130 oscillations min<sup>-1</sup>) at 30 °C. Ethanol was determined as described above.

**Dry weight determination.** The dry weights of suspensions of *M. trichosporium* OB3b were determined by heating 1 ml samples at 110 °C to constant weight.

**Protein determination.** Protein content of cell-free extracts was determined using the modified Lowry assay of Peterson *et al.* (1977). Bovine serum albumin was used as a standard.

**SDS-PAGE.** Electrophoresis was carried out using 12% (w/v) acrylamide gels using the discontinuous buffer system of Laemmli (1970). Proteins were fixed, stained with Coomassie blue and the gels destained as described by Laemmli (1970). The following were used as molecular weight standards: lactalbumin (mol. wt 14200); trypsin inhibitor, soybean (mol. wt 20100); trypsinogen, PMSF treated (mol. wt 24000); carbonic anhydrase from bovine erythrocytes (mol. wt 29000); glyceraldehyde-3-phosphate dehydrogenase, rabbit muscle (mol. wt 36000); albumin, egg (mol. wt 45000); albumin, bovine (mol. wt 66000). Methanol dehydrogenase of *M. trichosporium* OB3b (subunit mol. wt 60000) was used as a further standard. This was a gift from Dr D. J. Best, Cranfield Institute of Technology.

**Chemicals.** All chemicals were obtained from BDH except the molecular weight markers which were obtained from Sigma.

## RESULTS

### *Effect of copper ions on the intracellular location of MMO*

When *M. trichosporium* OB3b was grown on the low copper medium at biomass densities of 1.6–1.8 mg dry wt ml<sup>-1</sup>, the MMO activity was located predominantly in the soluble fraction of

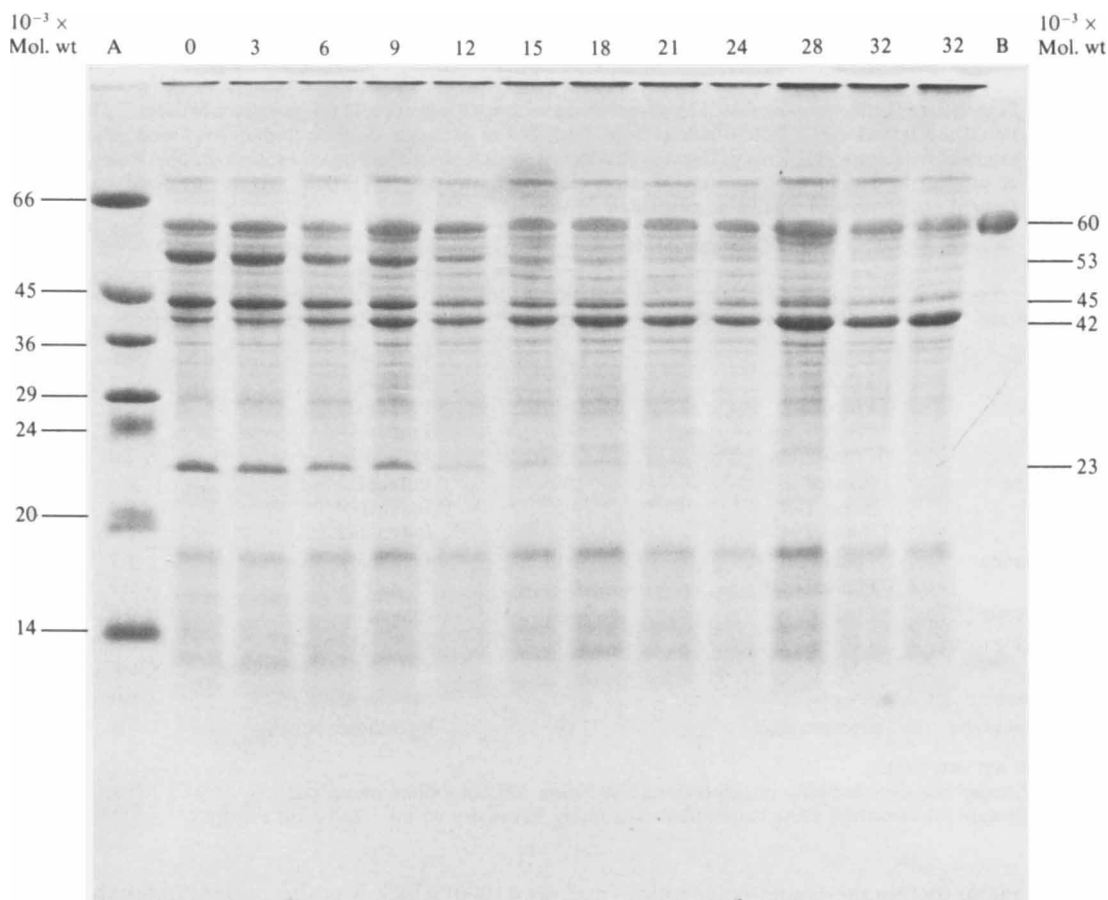


Fig. 1. SDS-PAGE of cell-free extracts from *M. trichosporium* OB3b during a switch from low to high copper medium. Cells were grown in continuous culture on low copper medium ( $1 \mu\text{M-Cu}^{2+}$ ; dilution rate,  $0.04 \text{ h}^{-1}$ ; biomass density,  $1.6\text{--}1.8 \text{ g dry wt l}^{-1}$ ). The concentration of  $\text{Cu}^{2+}$  was raised to  $5 \mu\text{M}$  by addition of  $\text{CuSO}_4$  solution to the chemostat and the feed was switched to high copper medium ( $5 \mu\text{M-Cu}^{2+}$ ). Samples were taken at intervals of 3–4 h for analysis of the cell-free extracts by SDS-PAGE. Electrophoresis was carried out as described in Methods. The numbers above each track indicate the time (h) after the switch to high copper medium. Track A shows the molecular weight markers (mol. wt 66000, 45000, 36000, 29000, 24000, 20100 and 14200) and track B is purified methanol dehydrogenase from *M. trichosporium* OB3b (subunit mol. wt 60000).

cell extracts [soluble activity was  $52 \text{ nmol min}^{-1} (\text{mg protein})^{-1}$ , particulate activity  $8.4 \text{ nmol min}^{-1} (\text{mg protein})^{-1}$ ]. By contrast, organisms grown on the high copper medium to the same biomass densities contained predominantly particulate MMO [soluble activity was  $4.8 \text{ nmol min}^{-1} (\text{mg protein})^{-1}$ , particulate activity  $62.5 \text{ nmol min}^{-1} (\text{mg protein})^{-1}$ ]. Organisms containing 100% particulate MMO were obtained by growing them on the high copper medium at a lower biomass density ( $1.0\text{--}1.1 \text{ mg dry wt ml}^{-1}$ ); this was accomplished by increasing the dilution rate of the cultures from  $0.04 \text{ h}^{-1}$  to  $0.043 \text{ h}^{-1}$ .

Differences in the amounts of major polypeptides were observed when the protein profiles of organisms containing either soluble or particulate MMO were analysed using SDS-PAGE. Three polypeptides (mol. wt 53000, 45000 and 23000) were present at high concentrations in the soluble fraction of organisms grown on the low copper medium (i.e. those containing soluble MMO) (Fig. 1). All three polypeptides were virtually absent from soluble fractions of organisms grown on the high copper medium which contained particulate MMO. These contained instead

Table 1. *Oxidation of n-alkanes, and aromatic and alicyclic compounds by suspensions of M. trichosporium OB3b containing either soluble or particulate MMO*

Suspensions (1 ml) were incubated in sealed flasks with each substrate (2 ml gaseous substrates, 2  $\mu$ l liquid substrates) over a time course (1 h) as described in Methods. Sodium formate was used as a source of reducing equivalents in all cases. Products were determined using gas chromatography. Rates of product formation are expressed as nmol min<sup>-1</sup> (mg dry wt)<sup>-1</sup>. Hexane was not oxidized by suspensions containing soluble or particulate MMO.

Substrate	Soluble MMO		Particulate MMO	
	Products detected	Initial rate of formation	Products detected	Initial rate of formation
Ethane	Ethanol*	ND	Ethanol*	ND
	Ethanal	25.0	Ethanal	35.0
Propane	Propanal	3.2	Propanal	2.4
	Propan-1-ol	6.2	Propan-1-ol	0.0
	Propan-2-ol	4.9	Propan-2-ol	12.6
Butane	Butanal	2.6	Butanal	0.6
	Butan-1-ol	4.2	Butan-1-ol	0.0
	Butan-2-ol	7.3	Butan-2-ol	5.9
Pentane	Pentanal†	1.1	Pentanal†	1.1
	Pentan-2-ol†	0.3	Pentan-2-ol†	0.5
Benzene	Phenol	74.0	Phenol	0.0
Ethylbenzene	1-Phenylethanol	5.7	1-Phenylethanol	0.0
	4-Hydroxyethylbenzene	13.0	4-Hydroxyethylbenzene	0.0
Styrene	Styrene oxide	82.0	Styrene oxide	0.0
Cyclohexane	Cyclohexanol	25.0	Cyclohexanol	0.0

ND, not determined.

\* Ethanol was detected only in suspensions containing 200 mM-sodium phosphate.

† Pentane was oxidized using suspensions containing 30 mg dry wt ml<sup>-1</sup> and 10  $\mu$ l substrate.

one major membrane-bound polypeptide (mol. wt 42000) which was also present in organisms grown on the low copper medium, although in much lower amounts (Fig. 1).

Particulate MMO activity was lost when membranes prepared from organisms grown on the high copper medium were solubilised using Triton X-100 (0.5%, v/v) or sodium deoxycholate (0.5%, w/v).

#### *Substrate specificities of soluble and particulate MMO*

*Oxidation of n-alkanes.* Table 1 lists the products resulting from the oxidation of *n*-alkanes by suspensions of *M. trichosporium* OB3b which contained either soluble or particulate MMO, and gives the initial rates of their formation. The figures given for primary alcohols cannot be regarded as true rates, since these are oxidized further to the corresponding aldehydes by the non-specific methanol (primary alcohol) dehydrogenase (Patel *et al.*, 1972). The aldehydes may also be oxidized further by a general aldehyde dehydrogenase to give the corresponding carboxylic acids (Patel *et al.*, 1980). Suspensions containing either soluble or particulate MMO oxidized ethane to acetaldehyde without accumulating ethanol. Formation of ethanol was observed, however, when suspensions were prepared in 200 mM-sodium phosphate buffer, showing that ethanol is the initial oxidation product. The accumulation of ethanol under these circumstances is due to inhibition of the methanol oxidation by high concentrations of phosphate (Higgins & Quayle, 1970), though there is no evidence that this is due to direct inhibition of methanol dehydrogenase. On raising the phosphate concentration from 20 mM to 200 mM there was a 91% decrease in the rate of ethanol oxidation by organisms containing particulate MMO, whilst MMO activity, measured by following the epoxidation of propene in the presence of 20 mM-sodium formate, decreased by only 25–30%. Suspensions of *M. trichosporium* OB3b containing soluble MMO catalysed both terminal and subterminal hydroxylation of propane and butane, as evidenced by the formation of the corresponding primary and secondary alcohols. Suspensions

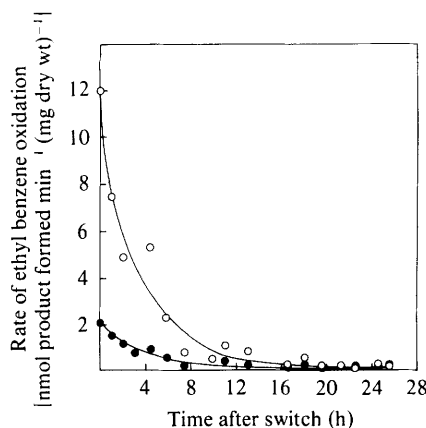


Fig. 2. Oxidation of ethyl benzene by *M. trichosporium* OB3b during a switch from low to high copper medium. Cells were grown in continuous culture on low copper medium ( $1\ \mu\text{M-Cu}^{2+}$ ; dilution rate,  $0.04\ \text{h}^{-1}$ ; biomass density,  $1.6\text{--}1.8\ \text{g dry wt l}^{-1}$ ). The concentration of  $\text{Cu}^{2+}$  was raised to  $10\ \mu\text{M}$  by addition of  $\text{CuSO}_4$  solution to the chemostat and the feed was switched to high copper medium ( $10\ \mu\text{M-Cu}^{2+}$ ). Samples ( $10\ \text{ml}$ ) were taken at intervals for determination of whole cell ethyl benzene oxidation rates. Organisms were incubated with shaking at  $30\ ^\circ\text{C}$  in  $250\ \text{ml}$  conical flasks in the presence of ethyl benzene ( $10\ \mu\text{l}$ ) and sodium formate ( $20\ \text{mM}$ ) as a source of reducing equivalents. Products, 4-hydroxyethylbenzene (○) and 1-phenylethanol (●), were analysed as described in Methods.

containing particulate MMO oxidized propane and butane to the corresponding secondary alcohols, but primary alcohols were not detected. Propanal and butanal accumulated, however, showing that terminal hydroxylation was occurring but that both the respective primary alcohols were oxidized further. Both MMOs oxidized pentane to pentanal (via pentan-1-ol) and pentan-2-ol, but oxidation at position 3 of the carbon chain was not observed. Oxidation products were not detected when suspensions or cell-free extracts were incubated with *n*-hexane. This was true whether they contained soluble or particulate MMO.

#### Oxidation of aromatic and alicyclic compounds

Organisms containing soluble MMO oxidized benzene, ethylbenzene, styrene and cyclohexane, yielding the products listed in Table 1. By contrast, none of these compounds were oxidised by organisms which contained only the particulate enzyme. This suggests that the two MMOs differ significantly in their capacity to hydroxylate cyclic hydrocarbons. Cultures grown on the low copper medium progressively lost the capacity to oxidize ethylbenzene when the concentration of cupric ions was raised from  $1\ \mu\text{M}$  to  $10\ \mu\text{M}$  (Fig. 2), yet their ability to oxidize propene remained constant [ $150\text{--}160\ \text{nmol min}^{-1}\ (\text{mg dry wt})^{-1}$ ]. These observations confirmed that the copper-induced transition from soluble to particulate MMO is accompanied by the loss of the ability to hydroxylate aromatic substrates. When the copper concentration of the fermenter feedstock was returned to  $1\ \mu\text{M}$ , the organism regained the ability to oxidize ethylbenzene (not shown).

#### DISCUSSION

This work corroborates the finding of Stanley *et al.* (1983) that copper availability determines the intracellular location (soluble or particulate) of MMO in *M. trichosporium* OB3b. This is also the case for *Methylococcus capsulatus* (Bath) (Stanley *et al.*, 1983). Research to date (Scott *et al.*, 1981a; Stanley *et al.*, 1983) has indicated that there are differences between soluble and particulate MMOs besides their intracellular location, which suggest that the particulate enzymes are not derived from the soluble enzymes merely by membrane attachment. Our results support and extend previous findings.

In both *M. trichosporium* OB3b and *M. capsulatus* (Bath) the inhibitor profile of the soluble MMOs differs considerably from that of the particulate enzyme (Scott *et al.*, 1981a; Stanley *et*

*al.*, 1983), and although both types of MMO can use NADH as an electron donor *in vitro*, Tonge *et al.* (1975) have suggested that *in vivo* the particulate enzymes may receive reducing equivalents from the electron transport chain or from NADH-independent dehydrogenases such as methanol dehydrogenase. This is supported by the recent observations of Leak & Dalton, (1983). Our findings show that the particulate MMO of *M. trichosporium* OB3b differs from the soluble enzyme in that it is unable to oxidize aromatic or alicyclic compounds (both enzymes oxidize methane, propene and *n*-alkanes). This suggests that the hydroxylase components of the two enzymes are different, and provides a further important distinction between the two types of MMO. It is of interest to note that the substrate specificity of the particulate MMO of *Methylomonas methanica* (Stirling *et al.*, 1979) is the same as that reported here for the corresponding enzyme from *M. trichosporium* OB3b. The ability to oxidize *n*-alkanes and *n*-alkenes but not aromatics or alicyclics may prove to be a general property of particulate MMOs, in which case it might be possible to determine whether or not organisms possess the soluble enzyme (which oxidize all four classes of compound) simply by measuring their capacity to oxidize a cyclic hydrocarbon such as ethylbenzene. This is clearly possible for *M. trichosporium* OB3b.

It has been suggested that methanotrophs might be used as biocatalysts in order to carry out epoxidations and hydroxylations on an industrial scale (Higgins *et al.*, 1981; Whittenbury & Dalton, 1982; Hou, 1984). Our findings shown that if *M. trichosporium* OB3b is to be used to oxidize cyclic hydrocarbons in an industrial process, then it will be essential to ensure that it is grown under conditions which yield soluble MMO. This restriction may well apply to other methanotrophs.

The soluble MMO of *M. capsulatus* (Bath) is now known to be a three-component enzyme (Colby & Dalton, 1976; Woodland & Dalton, 1984), and the corresponding enzyme from *M. trichosporium* OB3b seems to be essentially similar (Stirling & Dalton, 1979). By contrast, nothing is known of the polypeptide composition of the respective particulate enzymes. Stanley *et al.* (1983) have shown that three major polypeptides are present in cell-free extracts prepared from suspensions of *M. capsulatus* (Bath) which contain soluble MMO. These correspond to the three subunits (mol. wt 54000, 42000, 17000) of component A of the soluble MMO of this organism, and may constitute up to 30% of the total soluble protein (Woodland & Dalton, 1984). All three polypeptides disappear during the copper-induced transition from soluble to particulate MMO, whilst additional membrane-bound polypeptides accumulate (Stanley *et al.*, 1983). The levels of some of the major cellular polypeptides of *M. trichosporium* OB3b appear to change in a similar fashion, although it is not known at present whether any of these correspond to components of the soluble or particulate MMOs. This will only be established when both enzymes from *M. trichosporium* OB3b have been purified, and this is currently under investigation. This is likely to prove difficult for the membrane-bound enzyme, however, since attempts to solubilize the particulate MMOs of both *M. methanica* (Colby *et al.*, 1975) and *M. trichosporium* OB3b (the present work) have resulted in loss of activity.

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