

P-450 HEMEPROTEINS OF RHIZOBIUM JAPONICUM
Purification by Affinity Chromatography and
Relationship to P-450_{CAM} and P-450_{LM-2}*

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Summary: Rhizobium hemeproteins P-450a, b, and c cross react with antibodies to P-450_{CAM} and P-450_{LM-2}. Anti P-450_{CAM} IgG and phenobarbital, each bound to Sepharose 4B, were effective in purification of Rhizobium P-450c; the latter was more convenient. The amino acid composition of highly purified Rhizobium P-450c resembles the compositions of P-450_{CAM} and P-450_{LM-2}. These results suggest that P-450 heme proteins of unrelated substrate specificities may nevertheless contain similar structural features.

P-450 hemeproteins are usually not found in N₂-fixing organisms. The unexpected appearance of a family of soluble P-450's in both symbiotic Rhizobia (1,2) and cells grown anaerobically in a nitrate-organic carbon medium (3) therefore requires an explanation. Involvement of these cytochromes in an oxidative phosphorylation pathway which supports N₂-fixation in R. japonicum has recently been suggested (4) but their exact function remains unclear.

The wide distribution of P-450 hemeproteins in nature and their participation in seemingly unrelated metabolic processes has prompted us to raise the question of structural similarity between these b-type cytochromes. It is already known that P-450_{CAM} of the 5-exo hydroxylase of D (+) camphor of P. putida and P-450_{LM-2} of the phenobarbital-induced drug hydroxylase of rabbit liver microsomes resemble each other as judged from immunochemical and compositional data (5). The present communication extends this comparison of related P-450 hemeproteins to Rhizobial proteins. A first account of this work has been reported recently (6).

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Materials and Methods

Hemeproteins: Symbiotic N_2 -fixing cells of Rhizobium japonicum, strain CC705 (syn. Wisconsin 505), were isolated from "Lincoln" strain soybean root nodules (1). Partially purified Rhizobium hemeproteins P-450_a, b, and c were obtained from cell extracts by ion exchange chromatography on DE-52 cellulose (2). Partial purification of P-450_b which is found in non-symbiotic cells of Rhizobium was carried out using extracts of cells grown anaerobically in pure culture in the presence of nitrate (2,3). Crystalline P-450_{CAM} of P. putida (7) was generously provided by Dr. I. C. Gunsalus, and phenobarbital-induced liver microsomal P-450_{LM-2} containing 15 nmoles P-450 per mg protein (8) was kindly donated by Dr. M. J. Coon.

Antibodies and Radioimmunoassays: Rabbit antibodies to P-450_{CAM} and P-450_{LM-2} were obtained and purified as described previously (5). For competitive binding assays P-450_{CAM} and P-450_{LM-2} were labeled with ¹²⁵I exclusively at tyrosines (9) and without apparent loss of catalytic activity (10).

Affinity Chromatography: ω -Aminohexyl Sepharose 4B and ω -carboxypentyl Sepharose 4B were purchased from Pharmacia Fine Chemicals AB, Uppsala, Sweden. m-Aminophenobarbital was synthesized (11) and bound either to Sepharose 4B via BRCN coupling or to ω -carboxypentyl Sepharose 4B by carbodiimide dehydration (12). The capacity of these Sepharose derivatives was estimated to be 15-18 μ moles/ml for cytochrome P-450_{CAM}. Routinely, volumes of 0.25-0.35 ml of swollen resin were transferred into Pasteur pipettes, equilibrated with 0.05 M phosphate, pH=7.0, and immediately used for affinity chromatography of 3-5 mg of P-450 hemeprotein.

Isoelectric Focusing in Acrylamide Gel: The apparatus, constructed from Lucite to accommodate a gel bed of 25x15x0.5 cm, was cooled at 4° C. Filter paper strips (Whatman No. 3, 3x9 mm) soaked with sample solution to contain ~ 5 μ g protein were placed on the gel ~ 1 cm from the cathode. Electrofocusing was carried out in acrylamide gel (7% cross linkage, 2% ampholytes pH 3-10) for 2 h maintaining a constant current of < 35 mA. After the run the protein was fixed in 7% TCA.

Results and Discussion

Immunochemical Experiments: As seen in Fig. 1, the family of symbiotic Rhizobium P-450 hemeproteins showed substantial cross reactivity with anti P-450_{CAM} antibodies, namely, 54% for P-450_c, 50% for P-450_b, and 45% for P-450_a. This compares to 60% cross reactivity of P-450_{LM-2} with anti P-450_{CAM} antibodies reported earlier (5). It is deemed significant that the extent of displacement of Rhizobium hemeproteins by P-450_{CAM} is not the same for all members and that P-450_c shows the closest relationship to P-450_{CAM} in this assay. These immunochemical data are in agreement with a closer correlation of P-450_c and P-450_{CAM} with regard to absorption maxima (2), good solubility in aqueous buffers, and fairly acidic isoelectric points, at pH=4.9 and 4.5 for P-450_c and P-450_{CAM}, respectively (13). P-450_c also happens to be the most abundant

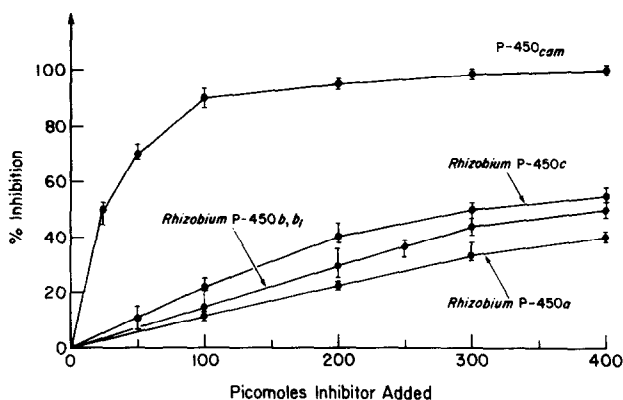


Fig. 1. Cross Reactivity of Rhizobium P-450 Hemeproteins with Antibodies to P-450_{CAM}.

Radioimmunoassays of competition of unlabeled Rhizobium P-450 hemeproteins with ¹²⁵I-labeled P-450_{CAM} for binding to anti P-450_{CAM} antibodies were performed in triplicate using 0.15 M NaCl-0.05 M sodium phosphate buffer, pH=7.0, as described previously (3). Measurements of radioactivity in the resulting precipitates carried out either with the ¹⁴C window of a Beckman Model LS-30 Liquid Scintillation Counter or with a Packard Tricarb Liquid Scintillation Spectrometer 3375, with β - γ -control, yielded identical sets of curves. The inhibition of binding by Rhizobium P-450 hemeproteins is based on maximal interaction of unlabeled P-450_{CAM} to anti P-450_{CAM} antibodies; these values were normalized to 100%.

Rhizobium P-450 component and the one most readily purifiable by ion exchange chromatography (2). P-450_{b₁} of anaerobic-nitrate grown cells, showing elution patterns in DE-52 ion exchange chromatography identical to that of symbiotic P-450_b (2), could not be distinguished from P-450_b by cross reactivity with anti P-450_{CAM} antibodies. Preliminary data not included here also point to strong cross reaction between Rhizobium P-450 hemeproteins a, b, and c and anti P-450_{LM-2} antibodies. In this case P-450_a seems to be the component most closely related to P-450_{LM-2}.

As outlined in Table I, the similarity in antigenic site structures was exploited for further purification of P-450_c via specific complexing with anti P-450_{CAM} IgG antibodies which had been immobilized on Sepharose 4B. In this case the batch procedure was preferred over column chromatography because of the need to incubate the P-450_c with the immobilized antibodies for several hours to guarantee complete interaction. Although this immunochemi-

Table I
Purification of Rhizobium P-450_{CAM} Using
Immobilized Anti-P-450_{CAM} Antibodies

- A. Preparation of Specific Antibodies:
- 1.) binding of P-450_{CAM} to Sepharose 4B via BrCN coupling
 - 2.) application of IgG fraction
 - 3.) removal of non-specific antibodies by washing
with 2 volumes of 50 mM $\text{PO}_4^{=}$ at pH 7.0
 - 4.) elution of 2 specific antibody fractions
 - a) 1 M NaCl, 50 mM $\text{PO}_4^{=}$, pH 7.0
 - b) 1 M NaCl, 10% dioxane, 50 mM $\text{PO}_4^{=}$, pH 7.0
 - 5.) dialysis of combined fractions; 80% recovery of anti-P-450_{CAM} antibodies.
- B. Batch Purification of P-450_C:
- 1.) binding of anti-P-450_{CAM} IgG to Sepharose 4B via BrCN coupling
 - 2.) incubation with Rhizobium P-450_C ($A_{280}/A_{417} = 1.02$): 18 hrs. at 4° C
 - 3.) removal of unbound protein contaminants by washing with 50 mM $\text{PO}_4^{=}$,
pH 7.0
 - 4.) elution of P-450_C with 1 M NaCl; 60% recovery of P-450_C ($A_{280}/A_{417} = 0.86$)

cal affinity procedure yielded P-450_C preparations of significantly increased purity as evidenced by the improvement of the spectral ratio, A_{280}/A_{417} , normally used as purity index, from 1.02 to 0.86 (Table I), several difficulties were encountered with this procedure. First, large amounts of pure P-450_{CAM} bound to Sepharose 4B via BrCN coupling were required for the preparation of specific antibodies, and second, only P-450_{CAM} antibodies of medium affinity toward P-450_C could be used effectively in the subsequent purification step. This selection was necessary to avoid losses due to loose binding of P-450_C or formation of complexes too tight to dissociate at moderately high salt concentrations.

Affinity Chromatography on Sepharose Derivatives: A more convenient and less costly purification procedure is summarized in Fig. 2. P-450_C bound only moderately strongly to ϵ -aminohexyl Sepharose 4B and was recovered in similar yields and even higher purity, $A_{280}/A_{417} = 0.74 - 0.78$, than from the immobilized antibody resin. m-Aminophenobarbital bound directly to Sepharose 4B via BrCN coupling was apparently not sufficiently exposed to permit good binding of P-450 hemeproteins but if it was attached to an extended alkyl arm

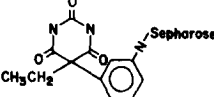
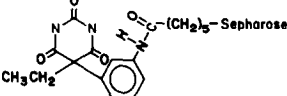
Column	P-450 a	P-450 b	P-450 c
ω -aminoethyl — H ₂ N-(CH ₂) ₆ -Sephacrose	+	++ A $\frac{280}{414}$ 3.1→2.6 73%	++ A $\frac{280}{418}$ 1.02→0.78 54%
Phenobarbital— 	+	+	+
Phenobarbital-alkyl=PBA 	+++ A $\frac{280}{416}$ 3.3→1.44 64%	+++ A $\frac{280}{414}$ 3.1→2.05 80%	+++ A $\frac{280}{418}$ 2.05→0.95 65%

Fig. 2. Affinity Chromatography of Symbiotic Rhizobium P-450.

Pasteur pipettes containing 0.25-0.35 ml of swollen Sepharose 4B derivatives were employed as miniature affinity columns. The resin was equilibrated with 0.05 M sodium phosphate, pH=7.0, and charged with 3-5 mg of P-450 dissolved in the same buffer. The columns were washed with 20 void volumes of buffer to remove contaminating proteins and the affinity-bound hemoproteins were then eluted with buffers of higher ionic strength: Weakly bound P-450s (+) eluted with 0.5 M phosphate, moderately strongly bound P-450s (++) eluted with 1.0 M phosphate-2.0 M NaCl, and very strongly bound P-450s (+++) eluted with 2.0 M urea, all at pH=7.0. All eluates were immediately dialyzed against 0.05 M phosphate.

(12) strong binding was observed for all Rhizobium P-450s as well as for P-450_{CAM} and P-450_{LM-2}. As shown by the improved spectral ratios, P-450s a, b, and c were substantially purified (Figs. 2+3). Better resolution of components was achieved by isoelectric focusing in polyacrylamide gel shown in Fig. 4. Each of the DE-52 fractions (2) was resolved into many bands including several hemoprotein bands with different isoelectric points. Thus we conclude that symbiotic Rhizobia contain at least 6 distinct P-450 hemoproteins. Affinity chromatography removed most of the non-heme contaminants (Fig. 4). Yet this preparation of P-450_c still contained two hemoproteins. Its spectral ratio, A₂₈₀/A₄₁₇ was 0.72-0.75. Since P-450_c contains fewer tyrosines than P-450_{CAM} or P-450_{LM-2} (see Table II), we estimate a hemoprotein content of about 70% for the purified preparation and assume that apo P-450_c contributes considerably to the remaining 30%.

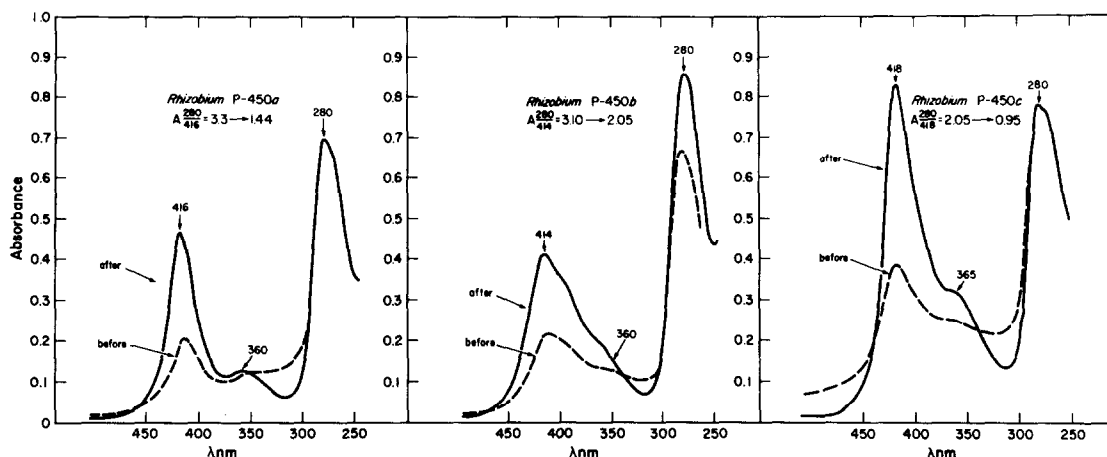


Fig. 3. Spectral Properties of Rhizobium P-450 Hemeproteins.

All absorption spectra were recorded with a Cary spectrophotometer, model 14, in 0.05 M phosphate, pH=7.0. The dashed and solid traces refer to the same protein preparations before and after affinity chromatography on PBA-Sepharose 4B respectively. The UV traces of P-450a and c before and after purification were matched but those of P-450b differed.

As expected, P-450_{LM-2} preparations (8) bound to α -phenobarbital alkyl (PBA) Sepharose very tightly and, after extensive washing with .5M NaCl in 50 mM phosphate pH 7.0, the hemeprotein eluted with 2 M NaCl in 50 mM phosphate pH 7.0 showed an improved spectral ratio, A_{280}/A_{417} of about 0.72 (about 95% pure).^{*} It was surprising, however, to find that all Rhizobium P-450's bound very tightly to PBA-Sepharose as well and had to be eluted with 1 M phosphate - 2 M NaCl. Even P-450_{CAM} bound quite tightly to this resin. It should be noted that none of these hemeproteins was retarded on Sepharose 4B or its - carboxypentyl or α -aminohexyl-p-aminobenzamidyl derivatives, and neither cytochrome c nor bovine serum albumin bound to PBA-Sepharose. Also, the P-450s did not bind at high salt concentrations as is preferred for sample

^{*}R. Goewert and K. Dus, unpublished data

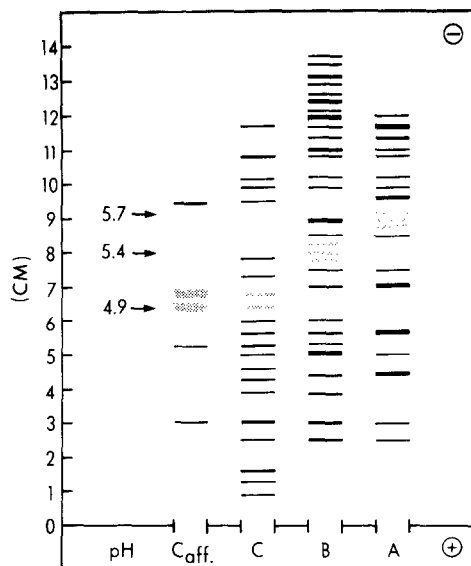


Fig. 4. Isoelectric Focusing of Rhizobium P-450 Hemeproteins in Polyacrylamide Gel.

Electrofocusing was carried out in gels of 7% cross linkage using 2% ampholyte solutions, pH 3-10, for 2h at 4° C maintaining a constant current below 35 mA. Rhizobium P-450 hemeproteins a, b, and c refer to fractions obtained from chromatography on DE-52 cellulose (2) while P-450 c_{aff} refers to P-450c purified by affinity chromatography on PBA-Sepharose.

application in hydrophobic chromatography (14). On the other hand, P-450s bound tightly to PBA-Sepharose exhibited the Soret maxima of the substrate-free forms rather than those of the substrate complexes.

The amino acid composition of the purified P-450 c preparation (Table II) is based on a molecular weight of 46,000 daltons as estimated by gel filtration on Sephadex G-75 (2). This composition shows close correlation to the compositions of P-450 $_{CAM}$ (13) and P-450 $_{LM-2}$ (5). All three P-450 hemeproteins are acidic and show an unusually high content of hydrophobic residues. The only significant deviations are the occurrence of only 2 cysteines and the much larger content of alanine in Rhizobium P-450 c . This protein seems to be more closely related to P-450 $_{CAM}$ than to P-450 $_{LM-2}$, but whenever it deviates from P-450 $_{CAM}$, as in the contents of Gly, Ile, Leu, and Arg, it tends to approach

TABLE II AMINO ACID COMPOSITION OF CYTOCHROMES

Amino Acids	P-450 _{CAM} <u>Pseudomonas putida</u>	P-450 _{LM-2} phenobarbital-induced rabbit liver microsomes	P-450 _C <u>Rhizobium japonicum</u>
Asx	36	35	33
Thr	19	23	14
Ser	21	26	20
Glx	55	43	54
Pro	27	24	25
CySH	6	6	2
Gly	26	30	33
Ala	34	23	54
Val	24	27	25
Met	9	8	8
Ile	24	19	14
Leu	40	46	46
Tyr	9	11	6
Phe	17	28	17
His	12	11	12
Lys	13	19	14
Trp	1	1	1
Arg	24	29	29
Total	397	409	407
Heme	1	1	1
P I	4.5	4.2	4.9

the values found for P-450_{LM-2}. Based on the resemblances among Rhizobium P-450_C, P-450_{CAM} and P-450_{LM-2} in immunochemical reactivities and amino acid compositions we suggest that these are structure-related P-450 hemeoproteins.

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