

Profile of Basic Carbon Pathways in Guard Cells and Other Leaf Cells of *Vicia faba* L.¹

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ABSTRACT

Guard cells and three other cell types from *Vicia faba* L. 'Longpod' leaflets were assayed for enzymes that catalyze one step in each of five major carbon pathways in green plants: the photosynthetic carbon reduction pathway (ribulose-bisphosphate carboxylase, EC 4.1.1.39), the photosynthetic carbon oxidation pathway (hydroxypyruvate reductase, EC 1.1.1.81), glycolysis ([NAD] glyceraldehyde-P dehydrogenase, EC 1.2.1.12), the oxidative pentose-P pathway (6-P-gluconate dehydrogenase, EC 1.1.1.44), and the tricarboxylic acid pathway (fumarase, EC 4.2.1.2). Neither ribulose-bisphosphate carboxylase nor hydroxypyruvate reductase could be detected in guard cells or epidermal cells; high levels of these activities were present in mesophyll cells. The specific activity of fumarase (protein basis) was about 4-fold higher in guard cells than in epidermal, palisade parenchyma or spongy parenchyma cells. (NAD) glyceraldehyde-P and 6-P-gluconate dehydrogenases also were present at high protein specific activities in guard cells (2- to 4-fold that in mesophyll cells).

It was concluded that the capacity for metabolite flux through the catabolic pathways is high in guard cells. In addition, other support is provided for the view that photoreduction of CO₂ by these guard cells is absent.

The distinct morphological differences and functions found when guard cells, ordinary epidermal cells, palisade parenchyma cells, and spongy parenchyma cells of C₃ dicotyledon leaves are compared (22) imply biochemical differentiation. Some of these biochemical differences are obvious, e.g. Chl is abundant in palisade parenchyma, but is generally lacking in ordinary epidermal cells. Other differences are more subtle; e.g. P-enolpyruvate carboxylase is elevated in guard cells (14). Investigators have attempted to determine differences by investigating enzyme distribution among various tissues, but few studies have limited the tissue sample to one particular cell type (*cf.* 14). Even these latter studies have been restricted to enzymes of the photosynthetic carbon reduction pathway and those involved in C₄ acid metabolism. To gain a better understanding of the overall biochemical functioning of these different cells, we have assayed them for enzymes that catalyze steps in major carbon pathways. We report the results in this paper.

MATERIALS AND METHODS

Materials. *Vicia faba* L. 'Longpod' were grown in a growth cabinet (600 μE m⁻² s⁻¹ at plant level, 60% RH; 14-h photoperiod:

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25°C/20°C). Fully expanded bifoliate leaves of 3 to 5-week-old plants were used. Analytical enzymes were from Boehringer; most other biochemicals were from Sigma.

Assays. Except for ribulose-bisP carboxylase (19), all assays were optimized with respect to substrate concentrations, buffer, and pH ("Specific Step," Table I) in 1-ml volumes with extracts of *Vicia* mesophyll protoplasts. NAD(P) oxidation/reduction was measured directly by fluorometry. The protoplasts were isolated as before (2) except that 10 mM ascorbate was included in the digestion medium. Extracts were made by rupture of the protoplasts by passage through a 15-μm screen. Chl was determined by the method of Arnon (1).

Pre-illuminated *Vicia* leaflets were quick frozen in melting N₂ and then freeze-dried at -40°C. Extracts of portions of these leaves were assayed for enzyme activity. Except for volume, the "Specific Step" and "First Indicator Step" were as given in Table I. (These controls were designed to verify enzyme activity stability and linearity with time and tissue concentration during steps equivalent to the initial ones used in the histochemistry experiments.) Other portions of these same leaves were used for dissection (40% RH, 19°C). Individual guard cell pairs, palisade parenchymal cells, spongy parenchyma cells, and pieces of epidermis (devoid of guard cells) were weighed on a quartz fiber balance (<20-ng samples). Purity of these samples has been documented earlier (4, 16). The assays were initiated by pushing the sample onto a "Specific Step" reagent droplet (Table I) using the oil well technique (9). In two assays (for ribulose-bisP carboxylase and fumarase), a "First Indicator Step" was required also. Subsequent steps were designed to measure NAD⁺ or NADPH by enzymic cycling (5, 7, respectively). Enzymic activities, calculated on a dry weight basis, were converted to a protein basis (20) or Chl basis (15, 21) or per cell basis (4, 16) using conversion factors previously published. General references to the micromethodology are References 7 and 13.

RESULTS

On a protein basis, the specific activity of fumarase was about 4-fold higher in guard cells than in the other cell types assayed (Fig. 1). On a Chl basis, fumarase-specific activity was 3- to 7-fold higher in guard cells than in the mesophyll (Table II). However, on a cell basis, fumarase-specific activity was lower in guard cells than in the mesophyll.

The specific activity (protein or Chl basis) of 6-P-gluconate dehydrogenase was higher in guard cells than in the epidermis or mesophyll (Fig. 1, Table I). On a weight basis, the specific activities were similar in all cell types (155 to 240 mmol/kg [dry weight]·h, Table II).

(NAD) glyceraldehyde-P dehydrogenase-specific activities were the highest of the enzymes assayed (Fig. 1, Table II). On a protein or Chl basis, the specific activity was highest in guard cells, although, on an absolute basis, there was more activity in mesophyll cells.

Table I. Analytical Flow Chart for Initial Steps of Enzyme Assays

Enzyme	Specific Step			First Indicator Step	
	Reagent	Blank/standard	Termination	Reagent	Termination
Fumarase (EC 4.2.1.2)	0.2 μ l 50 imidazole-Cl (pH 7), 10 fumarate; 0.02% (w/v) Triton X-100 for 60 min at 25°C	-Fumarate/ malate	pH shift by next reagent	0.2 μ l 250 2-amino-2- methylpropanol (pH 10.1), 0.4 NAD ⁺ , 20 glutamate; 2.5 μ g/ml malic dehydrogenase, 2.0 μ g/ml glutamate- oxaloacetate transami- nase for 30 min at 25°C	1 μ l 0.12 N NaOH followed by 80°C for 20 min
6-P-gluconate dehydrogenase (EC 1.1.1.44)	0.2 μ l 50 Tes (pH 7.5), 10 MgCl ₂ , 0.2 NADP ⁺ , 1 6-P- gluconate for 45 min at 25°C	-6-P-gluconate/ NADPH	1 μ l 0.12 N NaOH followed by 80°C for 20 min	None	
(NAD) glycer- aldehyde-P dehy- drogenase (EC 1.2.1.12)	0.2 μ l 100 Tris-Cl (pH 7.8), 10 MgCl ₂ , 10 2-mercaptoetha- nol, 0.5 ATP, 0.2 NADH, 2 3-P-glycerate; 0.5 μ g/ml di- alyzed P-glycerate kinase for 60 min at 25°C	-3-P-glycerate/ NAD ⁺	1 μ l 0.12 N HCl followed by 80°C for 20 min	None	
Hydroxypyruvate (glyoxylate) reductase (EC 1.1.1.81)	0.2 μ l 100 Mes (pH 6), 0.4 NADH, 250 glyoxylate; 0.01% (w/v) Triton X-100 for 60 min at 25°C	-Glyoxylate/ NAD ⁺	1 μ l 0.15 N HCl followed by 80°C for 20 min	None	
Ribulose-bisP carboxylase (EC 4.1.1.39)	0.1 μ l (or 0.5 μ l) 50 Tris-Cl (pH 8.1), 10 MgCl ₂ , 1 GSH, 20 NaHCO ₃ , 0.2 ribulose- bisP; 0.02% (w/v) BSA for 15 min at 25°C	-Ribulose-bisP/ 3-P-glycerate	pH shift by next reagent	0.1 μ l (or 0.5 μ l) 100 imid- azole Cl (pH 6.6), 0.03 NADH, 0.1 EDTA, 0.3 ATP, 20 NaCl; 100 μ g/ ml dialyzed (NAD) glyceraldehyde-P dehy- drogenase, 40 μ g/ml di- alyzed 3-P-glycerate ki- nase for 15 min at 25°C	1 μ l 0.3 N HCl followed by 80°C for 20 min

Neither ribulose-bisP carboxylase nor hydroxypyruvate reductase was detected in guard cells or epidermis (Fig. 1, Table II). There was no striking difference in the distribution of the specific activities of these two enzymes between the mesophyll cell types.

DISCUSSION

It was reported earlier that enzymes of the photosynthetic carbon reduction pathway are present in both mesophyll cell types (15, 19). These enzyme activities are insignificant in guard cells (8, 19, 23; W. H. Outlaw, Jr., M. C. Tarczynski, L. Anderson, unpublished) and ordinary epidermal cells (19) of C₃ plants. (There is immunological evidence, however, for ribulose-bisP carboxylase in guard cells of some Crassulacean acid metabolism plants [8]). Our results (Fig. 1, Table II) confirm these previous reports and are included here for purposes of comparison. Our inability to detect hydroxypyruvate reductase, a marker enzyme for the photosynthetic carbon oxidation pathway, in cells lacking the photosynthetic carbon reduction pathway supports the conventional view (6, 26) of the relationship between these two carbon pathways. Thus, it seems clear that both these pathways are absent in guard cells of C₃ plants. However, PSII is present (11, 21, 28,

but see 24).³ Because guard cells possess PSII and lack (NADP) glyceraldehyde-P dehydrogenase, the electron flow in guard cell chloroplasts may serve to sense PAR (21). There are data that show guard cells are sensitive to PAR (25, A. Schwartz and E. Zeiger, unpublished) and that red light and blue light act synergistically in stimulating malate production in epidermal peels (12). In addition, several enzyme activities extracted from epidermal peels are modulated by thiol reagents (L. M. Rao and L. Anderson, unpublished), and thiol-inhibiting reagents prevent stomatal opening (10). Conceivably, this modulation might be effected *in situ* by linear electron transport, but we are aware that these effects may be brought about by other mechanisms.

(NAD) glyceraldehyde-P dehydrogenase activity, 6-P-gluconate dehydrogenase activity, and fumarase activity were higher in guard cells than in the other leaf cells (protein or Chl basis). If the elevated activities of these enzymes are correlated with high potential fluxes through glycolysis, the oxidative pentose-P pathway, and the tricarboxylic acid cycle, these findings indicate that

³ Any lingering doubt that PSII is present in guard cells can be dismissed on the basis of our unpublished results (K. C. Vaughn and W. H. Outlaw, Jr.). We have demonstrated with electron microscope-level histochemistry (see 27) the presence of PSII in barley guard cells. We have also detected DCMU-sensitive variable fluorescence in individual guard cell pairs of broad bean using an ultramicrofluorometer.

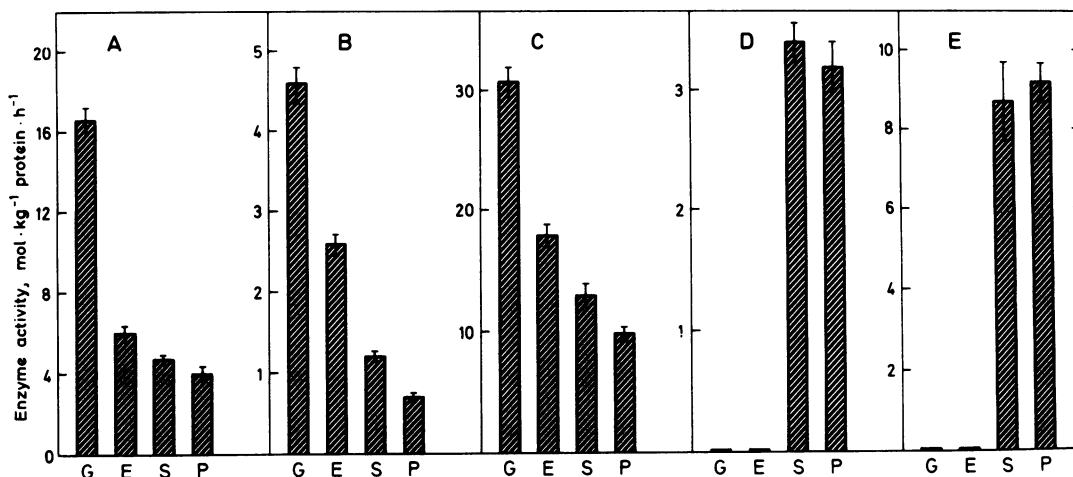


FIG. 1. Protein specific activity of fumarase (A), 6-P-gluconate dehydrogenase (B), (NAD) glyceraldehyde-P dehydrogenase (C), hydroxypyruvate reductase (D), and ribulose-bisP carboxylase (E), in guard cells (G), epidermal cells (E), spongy parenchyma cells (S), and palisade parenchyma cells (P) dissected from freeze-dried *V. faba* leaflet. Bars are SE. $n = 20$ (A-D) or $n = 6$ (E).

Table II. Enzyme Specific Activities in Guard Cells and Other Cells of *V. faba* L. Leaflet

Statistics are in Figure 1.

	Fumarase	6-P-Gluconate Dehydrogenase	(NAD) Glyceraldehyde-P Dehydrogenase	Hydroxypyruvate Reductase	Ribulose-bisP Carboxylase
Guard cells					
mmol/kg (dry wt)·h	860	240	1600	N.D. ^a	N.D.
μmol/mg Chl·h	576	160	1072	N.D.	N.D.
pmol/cell pair·h	5.2	1.4	9.6	N.D.	N.D.
Epidermal cells					
mmol/kg (dry wt)·h	440	190	1300	N.D.	N.D.
Spongy parenchyma cells					
mmol/kg (dry wt)·h	820	205	2300	610	1530
μmol/mg Chl·h	186	46	522	138	347
pmol/cell·h	11	2.8	32	8.4	21
Palisade parenchyma cells					
mmol/kg (dry wt)·h	850	155	2100	670	1940
μmol/mg Chl·h	80	15	199	64	184
pmol/cell·h	9.9	1	24	9	22

^a N.D., not detectable.

guard cells have evolved high capacity for heterotrophic energy-converting processes. This interpretation is consistent with the energy requirements for ion accumulation (e.g. 3), oxaloacetate reduction (e.g. 18), and starch catabolism (e.g. 17) during stomatal opening.

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