Isolation of Cytoplasmic Enzymes from Pollen¹

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ABSTRACT

The cytoplasmic isozyme of many cytoplasmic-organelle isozyme pairs, as well as other cytoplasmic enzymes in plants, can be readily obtained from pollen by soaking it in an appropriate buffer for 4 hours. Enzymes localized in subcellular organelles appear not to be released during the soaking period, although they are released if the pollen is crushed. The technique is a useful initial step in studies of subcellular localization of enzymes or for obtaining small quantities of cytoplasmic enzymes free of organellar contaminants.

The intracellular localization of enzymes has provided considerable information regarding the compartmentation of metabolic pathways. Techniques are now available for the isolation of various organelles from plant tissues (see ref. 19 for recent review). However, the cytoplasm and its complement of enzymes has been difficult to obtain free from contamination from organelle enzymes because techniques used to disrupt the plant cell also disrupt a certain proportion of the organelles. Cytoplasmic enzymes have been obtained following protoplast formation (14), but this procedure requires care and practice for optimal results. We report that many cytoplasmic enzymes can be obtained from pollen of many plant species by a simple and rapid method.

MATERIALS AND METHODS

Fresh pollen was collected from *Pisum sativum* L. cv. Alaska and *Spinacia oleracea* L. cv. Early Hybrid No. 7 because these species have been favorite materials for enzyme localization studies. In addition, we examined enzymes released from pollen of *Zea mays* L., *Clarkia unguiculata* Lindley, *Cedrus atlantica* Manetti, *Oenothera biennis* L., *Lycopersicon esculentum* Mill., and *Stephanomeria exigua* ssp. *coronaria* Gottlieb. The pollen was usually placed directly in buffer, although dry storage for 24 h at room temperature did not seriously impair the analysis. The buffer was generally 50 mM Tris-HCl, pH 8.0, containing 1 mM EDTA and 14 mM ME². Both 100 mM K-phosphate, pH 7.0, and 50 mM EDTA, pH 8.6, were also used. Approximately 5 mg pollen were placed in 0.5 ml chilled buffer and allowed to soak for 1 to 12 h. The extract then was used for starch gel electrophoresis. Crushed pollen extracts were obtained by grinding the soaked pollen suspension by hand in a chilled mortar for 1 to 2 min, which resulted in approximately 70% cell breakage. The slurry was centrifuged for 2 min at 7,000g. The supernatant fraction was removed by pipette and the pellet was discarded. Extracts from green leaf tissue were obtained by grinding 0.5 g tissue in 0.5 ml chilled 50 mm K-phosphate, pH 8.2, containing 2 mm EDTA and 14 mm ME.

Starch Gel Electrophoresis. Starch gels were prepared and electrophoresis was conducted as described previously (7). Assays were as follows: PGI (23); PGM (modified from (4) to contain 3 mM disodium α -D-glucose-1-P and 0.9 μ M dipotassium α -D-glucose-1,6-diP); AAT (7); ADH (21); ALDO (modified from ref. 7 by using 100 mM Tris-HCl, pH 8.5, as the buffer and 0.3 mM MTT in place of NBT); NAD- and NADP-specific GA3PDH (with 0.13 mM NADP in place of the NAD when assaying for the NADPspecific form) (4); catalase (23); TPI (modified from ref. 28 by using 100 mM Tris-HCl, pH 8.2, as the buffer and including 0.3 mM MTT and 100 μ M PMS); NAD-specific MDH (100 mM malate-Tris, pH 7.2, containing 0.4 mM MTT, 100 μ M PMS, and 0.45 mM NAD); SKDH (100 mM Tris-HCl, pH 8.5, containing 1.5 mM shikimic acid, 0.3 mM MTT, and 100 μ M PMS).

Time-Course Studies and Spectrophotometric Assays. Approximately 10 mg *Clarkia* pollen, weighed to the nearest 0.1 mg, was placed in an Eppendorf microcentrifuge tube containing 0.5 ml of 50 mM K-phosphate, pH 8.5, with 1 mM EDTA and 14 mM ME. The mixture was gently shaken by hand and allowed to stand at room temperature. Periodically, 0.025-ml aliquots were taken for enzyme assays at 22 C on a Zeiss PM6 recording spectrophotometer using a 0.5-ml cuvette. Catalase activity was measured using the method of Lück (15); fumarase, by the method of Racker (20). The assay mixtures for PGI, NAD- and NADP-specific GA3PDH, ALDO, ADH, and MDH were the same as those given above except that MTT (or NBT) and PMS were omitted. For these enzymes, the activity was determined by measuring the production of reduced dinucleotides at 340 nm. Blanks lacking substrate were run to allow correction for interfering reactions.

Chloroplast Isolation. Plants used for chloroplast extractions were kept in the dark for 16 h prior to harvest in order to reduce chloroplast starch content. Pea chloroplasts were extracted in a buffer modified (from ref. 17) to contain 0.33 M sorbitol, 100 mM Tricine (pH 7.5), 10 mM KCl, 1.5% (w/v) Ficoll, 0.1% (w/v) BSA, and 1 mm isoascorbic acid. Five g leaf tissue were sectioned into 1- to 2-cm² pieces and homogenized in 30 ml chilled extraction buffer by five 1-s bursts on a Sorvall Omni-Mixer. The slurry was filtered through four layers of Miracloth, with glass wool inserted between layers, and then centrifuged for 60 s at 1000g. The supernatant was discarded, and the pellet was resuspended in 20 ml extraction buffer and centrifuged again for 60 s at 1000g. Spinach chloroplasts were extracted in the buffer described in reference 3 except that 25 mm Hepes, pH 7.5, was used instead of Mes, and sodium pyrophosphate was not added. The initial homogenization and centrifugation steps were similar to those used for the pea preparation.

The chloroplast pellets of both species were further purified in

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² Abbreviations: ME, 2-mercaptoethanol; PGI, phosphoglucose isomerase; PGM, phosphoglucomutase; AAT, aspartate aminotransferase; ALDO, fructose-1,6-diphosphate aldolase; GA3PDH, glyceraldehyde-3phosphate dehydrogenase; TPI, triose phosphate isomerase; MDH, malate dehydrogenase; SKDH, shikimic dehydrogenase; MTT, 3-(4,5-dimethylthiazolyl-2)-2,5-diphenyltetrazolium bromide; NBT, nitro blue tetrazolium; PMS, phenazine methosulfate.

the following manner. Each was resuspended in 6 ml of the extraction buffer initially used for each species and placed on top of a 20 to 55% continuous sucrose gradient. This was centrifuged for 5 min at 4000 rpm and then for 5 min at 7000 rpm in a Sorvall HS-4 swinging bucket rotor. Two discrete Chl-containing bands were present after centrifugation. The lower band, containing the intact chloroplasts, was collected and divided into two aliquots. One of these was used for electron microscopy, and the other was diluted with an equal volume of 50 mM phosphate buffer (pH 8.2), containing 2 mM EDTA and 14 mM ME, and centrifuged for 2 min at 7000g. The supernatant fraction was used as the chloroplast extract in the electrophoretic studies.

Buffer Concentration Study. To test the effect of increasing the osmotic potential of the soaking buffer, approximately 10 mg *Clarkia* pollen was placed in 0.5 ml of each of three solutions: (a) 50 mM K-phosphate, pH 8.2, containing 2 mM EDTA and 14 mM ME; (b) the above solution supplemented with 1.0 M mannitol; (c) the chloroplast extraction buffer (without the isoascorbic acid but including 14 mM ME). The mixtures were shaken gently and allowed to stand at room temperature. After 30, 60, 120, and 270 min, 0.025-ml aliquots were taken from each solution, and the PGI activity was measured on the spectrophotometer. In a parallel study, 2.5 mg pea pollen was placed in 0.5 ml solution (a) above, and a second 2.5 mg was placed in 0.5 ml of the same solution containing 1% (w/v) NaCl since this has been reported to inhibit germination and prevent the pollen membrane from bursting (16). Aliquots of 0.025 ml were taken at 30 and 120 min for PGI assays.

RESULTS

The electrophoretic patterns of the various enzyme systems were examined in extracts of whole leaf, soaked pollen, and whole chloroplast preparations. The purities of the spinach and pea chloroplast preparations were verified by electron microscopy.

PGI, TPI, and PGM were examined by electrophoresis in both spinach and pea. For each system, the leaf extracts of both species contained two isozymes (Table I). The spinach chloroplast extract contained only the more anodal isozymes of PGI, TPI, and PGM. These results confirm previous observations that the more anodal isozymes of PGI (22) and PGM (18) are plastid-specific. The spinach soaked-pollen extract contained only the less anodal isozyme of these three systems. The pea chloroplast extract contained only the more anodal PGI and the less anodal TPI and PGM isozymes. The pea soaked-pollen extract contained only the less anodal PGI and the more anodal TPI and PGM (Fig. 1). For both species, the chloroplast isozyme of each system was present within pollen and could be obtained as an additional isozyme by soaking the pollen for a prolonged time or by mechanical crushing (Fig. 2). Thus, the plastid isozymes are present but retained within the pollen, whereas the cytoplasmic isozymes are readily released.

Two AAT isozymes could be clearly resolved by electrophoresis in pea leaf extracts. A third AAT isozyme with intermediate mobility was occasionally observed. The least anodal isozyme was present only in soaked-pollen extracts, and the most anodal one was present only in the chloroplast extract (Table I).

The most anodal MDH isozyme of pea, previously reported to be in the cytoplasm (29), was found only in the soaked-pollen extracts. None of the MDHs was found in the chloroplast extracts. Their compartmentation was not studied further. Electrophoresis revealed only a single enzyme for each of the other systems examined in pea. The plastid marker, NADP-specific GA3PDH, was present in extracts from whole leaves and the chloroplasts, but not in soaked pollen. However, the enzyme was present in the crushed-pollen extracts. Similar results were obtained for ALDO and SKDH (Table I).

The peroxisomal marker catalase exhibited low activity in soaked pollen extracts, suggesting that it is not entirely confined to extracytoplasmic compartments in pollen. Indeed, cytoplasmic catalase has previously been reported in castor bean (6). Catalase activity was 20-fold higher in the crushed-pollen extracts.

The mitochondrial marker fumarase was not resolved by electrophoresis. However, spectrophotometric analysis indicated it was present in the crushed-pollen extracts but absent from soaked pollen.

The cytoplasmic marker NAD-specific GA3PDH was present in extracts from soaked pollen. Thus, all of these results are consistent with the model that only cytoplasmic enzymes are readily extracted from pollen.

A number of enzymes were also examined in soaked-pollen extracts of *C. unguiculata* in order to test the generality of this finding. The soaked pollen extracts contained the cytoplasmic enzymes NAD-specific GA3PDH, ADH, and MDH but did not contain the organelle markers NADP-specific GA3PDH and fumarase. The cytoplasmic isozyme of PGI was present in the pollen extract but not the more anodal chloroplast PGI. We also examined PGI in *O. biennis, L. esculentum,* and *S. exigua* ssp. coronaria. In all three species, two regions of PGI activity were observed after electrophoresis of green leaf extracts, the more anodal being localized in the chloroplast fraction (unpublished data). Soaked-pollen extracts from each species contained only the more cathodal PGI isozyme.

In contrast to the rapid release of cytoplasmic enzymes from the pollen of *Clarkia*, the soaked-pollen extracts of maize and *Cedrus* were unusual in that they contained very little activity of PGI or ADH (other enzymes were not tested). Freezing or thawing their pollen before soaking yielded higher activity, although this treatment also resulted in release of small amounts of the plastid isozymes.

The time-course experiment indicated that PGI, NAD-specific GA3PDH, MDH, and ADH all diffused from *Clarkia* pollen shortly after it was immersed in the buffer. The activities of these enzymes increased more or less at a linear rate for the first 90 min and were essentially completed after 4 h soaking (Fig. 3). Evidence that a significant portion of the cytoplasmic PGI was not released into the buffer, even after 24 h soaking, was obtained by transferring soaked pollen into fresh buffer and crushing the pollen. Both the plastid and the cytoplasmic forms of PGI were obtained after crushing. Since both isozymes were present in this extract, their relative activity could not be determined spectrophotometrically.

The effect of increasing the osmotic potential upon the release of PGI was also tested using *Clarkia* pollen. The presence of 0.33 M sorbitol or 1 M mannitol in the extraction buffer did not affect the amount or rate of enzyme release, evaluated as per cent of the activity released in the crushed-pollen extract (Table II). The total PGI activity in the crushed pollen was similar in the control and in both test buffers (9.5 \pm 2.2 units/mg pollen; one unit is defined as the amount of enzyme that causes an increase of 0.1 A units/ min under the above conditions.

In a parallel experiment, the influence of 1% NaCl upon PGI release from pea pollen was also tested. No difference was noted in the PGI activities in the two buffers (Table II). The total PGI activity in the crushed pea pollen was 4.7 ± 0.3 units/mg pollen; units are defined as above.

DISCUSSION

Nearly a century ago, Green (8) mentioned that moistened pollen grains released a substance, presumably amylase, that liquified starch paste. Our data indicate that only cytoplasmic enzymes are released from pollen during the first 12 h of soaking in buffer, whereas organelle-bound enzymes are not. Consequently, pollen provides a convenient source of these enzymes. That enzymes are readily released from pollen is well known and has been demonstrated in *Iris* (10), *Phalaris* (12), *Ambrosia* and *Gladiolus* (13), and *Salsola, Pyrus*, and *Plantago* (25), in addition to the species studied by us. Stanley and Linskens (24) present a

	Table I.	Isozymes I	Resolved by	Electrophore	sis of Leaf,	Chloroplast,	and Pollen	Extracts
Numbers	indicate	mobility of	isozyme rel	lative to the	borate from	t.		

	Species	Isozymes					
Enzyme		Leaf	Chloro- plast	Pollen	Intracellular Location	Reference	
PGI	Pea	0.51	0.51		Chloroplast	27	
		0.39		0.39	Cytoplasm	27	
	Spinach	0.47	0.47		Chloroplast	22	
		0.28		0.28	Cytoplasm	22	
TPI	Pea	0.54		0.54	Cytoplasm	1	
		0.48	0.48		Chloroplast	1	
	Spinach	0.54	0.54		Chloroplast	This paper	
		0.44		0.44	Cytoplasm	This paper	
PGM	Pea	0.66		0.66	Cytoplasm	27	
		0.58	0.58		Chloroplast	27	
	Spinach	0.56	0.56		Chloroplast	18	
		0.42		0.42	Cytoplasm	18	
AAT	Pea	0.37	0.37		Chloroplast	27	
		0.27		0.27	Cytoplasm	27	
MDH	Pea	0.48		0.48	Cytoplasm	29	
		0.45			Microbody	29	
		0.20			Mitochondria	29	
Aldolase	Pea	0.36	0.36		Chloroplast	2	
GA3PD(NAD)	Pea	0.25		+ª	Cytoplasm	19	
GA3PD(NADP)	Pea	0.18	0.18		Chloroplast	19	
Catalase	Pea	0.15	ND⁵	faint	Peroxisome	19	
SKDH	Pea	0.33	0.33		Chloroplast	5	

^a GA3PDH activity in soaked pollen extracts from pea was too low to produce a clear stain on starch gels but could be measured using the spectrophotometric assay.

^b ND, not determined.



FIG. 1. Electrophoretic patterns of isozymes of PGM, PGI, AAT, and TPI on starch gels. Comparison of chloroplast (C), leaf (L), and soaked-pollen (P) extracts from pea. Anode is at the top of the figure.

list of approximately 80 enzymes that have been detected in pollen extracts.

The absence of enzyme activity in a soaked-pollen extract does not necessarily indicate that the enzyme is localized within an organelle. The enzyme could be attached to the ER or other membrane, may not be expressed in pollen, or may simply not be leachable [e.g. maize and also pine and palm (25)]. The technique



FIG. 2. Electrophoretic patterns of PGI isozymes in spinach extracts. A, crushed-pollen extract; B, soaked-pollen extract; C, leaf extract. Anode is at the top of the figure.

is a worthwhile initial step in studies of subcellular localization and for obtaining small quantities of cytoplasmic enzymes free from organellar contaminants. Its simplicity recommends it.

Our attempts to determine the total activity of specific enzymes per mg pollen proved difficult because the amount of activity released from the pollen appears to be affected by its age, the age of the plant, and environmental conditions. In addition, the amount of enzyme may be species-specific; thus, *Clarkia* and pea pollen extracts had different amounts of PGI activity.

Previous studies (26) of the genetic basis for the cytoplasmic PGI in *Clarkia* demonstrate that this enzyme, at least, is synthesized by the pollen and not by surrounding tapetal cells. This follows because the intragenic hybrid enzyme characteristic of individuals heterozygous for the coding gene locus is not observed in soaked pollen extracts. If the PGI were of tapetal origin, all three forms of the enzyme (*i.e.* both homodimers and the heterodimer) would be expected in the pollen of such plants. Other studies (9, 10) showed that acid phosphatase, ribonuclease, and esterase that leach from pollen are also synthesized therein.



FIG. 3. Results of time-course study for PGI (O---O), MDH ---O), ADH (×-----×), and NAD-specific GA3PDH (+-----+) from Clarkia pollen. Values on ordinate are percentage of enzyme activity in extract relative to that found after 24 h soaking. The PGI data points are the average of three independent experiments. Data points for MDH, ADH, and NAD-specific GA3PDH are the average of two pollen extracts examined during a single experiment. Standard errors for the data points are similar to those given in Table II.

Table II. Effect of Buffer Solute Concentration on Leaching Process

PGI activity in soaked pollen extracts as a percentage of activity in the crushed pollen extract. Values are the means \pm sE of three independent extractions with two replicates each for Clarkia pollen in 0.05 M phosphate, and one experiment with two replicates for the sorbitol test and three replicates for the mannitol tests. The test in pea was a single experiment with two replications.

Time		Clarkia PG	Pea	Pea PGI		
after Immer- sion	0.05 м Phos- phate	0.33 м Sorbitol	1.0 м Mannitol	0.05 м Phos- phate	1% NaCl	
min			%			
30	6.0 ± 1.2	9.0 ± 5.0	10 ± 0.8	15 ± 1.0	13 ± 1.0	
60	16 ± 2.6	26 ± 2.0	25 ± 9.0			
120	42 ± 6.6	45 ± 1.5	48 ± 7.4	23 ± 0.2	20 ± 0.3	
270	60 ± 12	53	53 ± 7.8			

The mechanism of enzyme release from the pollen appears to be the same for all of the enzymes we assayed since they followed a similar time course. Their release even in buffers having high osmotic potential may mean that they are located in the intine layer as proposed for other enzymes (11, 12). Alternatively, the pollen plasmalemma may be more "porous" than the membranes surrounding the organelles. Regardless of the actual mechanism of enzyme release, pollen is likely to serve as a useful source of cytoplasmic enzymes.

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