Regulation by Phospholipids and Kinetic Studies of Plant Membrane-Bound UDP-Glucose Sterol β -D-Glucosyl Transferase'

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ABSTRACT

Solubilization and partial purification of the microsomal UDP-glucose sterol glucosyl transferase activity from maize coleoptiles by chromatography on DEAE-cellulose resulted in a highly delipidated (>95%) and inactive enzymic preparation. Addition of sterols revealed part of the activity and subsequent addition of phospholipids further increased the activity. Negatively charged phospholipids were shown to be by far the best activators. The purification step also produced the elimination of two interfering microsomal enzymic activities: UDPase and steryl glucoside acyl transferase. The removal of these two enzymic activities was a prerequisite for kinetic studies including product-inhibition studies, since the substrates of these two latter enzymes are the products of UDPG-SGTase activity. The results of the kinetic studies strongly suggest an ordered bi-bi mechanism for the glucosylation of sterols. Finally the effect of different phospholipids on the kinetic parameters of the reaction was studied. Both phosphatidylcholine and phosphatidylglycerol significantly decrease $K_{m\text{-stero}l}$ (and not $K_{m\text{-}UDPyhence}$) and increase the reaction V_{max} . The decrease of $K_{\text{m-sterol}}$ is similar with both phospholipids whereas the increase of V_{max} is much greater with phosphatidylglycerol than with phosphatidylcholine.

Steryl glucosides $(SG)^2$ and acylated steryl glucosides (ASG) are present with free and esterified sterols in all plant tissues investigated so far, most probably as components of membrane structures (30). The glucosylation of sterols is catalyzed by UDPG-SGTase, a plasma membrane bound enzyme of plant cells (1 1). The reaction consists of a glucose transfer from UDPglucose to a phytosterol with formation of a β -glucosidic linkage between the anomeric carbon of glucose and the 3-hydroxyl group of the sterol (30):

 $Sterol + UDP-glucose \rightarrow steryl glucose + UDP$

Sterols are major constituents of the plasma membrane and their role in the regulation of membrane fluidity is now well established (9). By controlling the amount of free sterols in the membrane, UDPG-SGTase might participate in the regulation of membrane fluidity.

In a recent work we showed the phospholipidic dependence of maize coleoptile UDPG-SGTase: partial delipidation of the enzymatic preparation by acetone precipitation (29), or selective solubilization (5), results in a strong inhibition of the enzymic activity; the addition of PL to the delipidated enzymic preparation fully restores the activity (5, 29).

The regulation of membrane-bound enzymes by PL is well established (for a review, see Refs. 23 and 24), but papers dealing with the lipid-dependence of plant membrane-bound enzymes are just a few (8, 12, 14-16, 20, 23, 28). Some enzymes exhibit a certain specificity in the lipid requirement for optimal activity: most of them belong to bacterial or animal systems (23, 24); among plant lipid-dependent enzymes, only one case was described (20).

Investigation of the lipid dependence of UDPG-SGTase on a molecular basis required purification and total delipidation of the enzyme as well as knowledge of the kinetic parameters of the enzymic reaction. In the present paper we report the partial purification of the solubilized UDPG-SGTase activity from etiolated maize coleoptiles microsomes. The purification-associated delipidation allowed a study of the dependence towards different PL. In addition, the kinetic mechanism of the purified enzyme could be analyzed and we studied the effect of various PL on the kinetic parameters of the reaction.

Preliminary data from some aspects of this work have been reported (6).

MATERIALS AND METHODS

Chemicals. Technical sitosterol from Fluka, Switzerland consisted in 93% β -sitosterol, the remainder being campesterol, according to GLC analysis (3). Sitosteryl glucoside was ^a generous gift of Professor Pegel, Durban, South Africa. UDP, UDP-glucose, 98 to 99% pure phospholipids, and free and esterified fatty acids were from Sigma. UDP-[U-'4CJglucose (240 mCi/mmol) was purchased from Amersham (U.K.) and Emulphogene BC 720, from Gaf Corp. (New York).

Plant Material Preparation of Membrane Fraction and Solubilization of UDPG-SGTase Activity. The cultivation of maize seedlings (5) and preparation of microsomes from coleoptiles (29) have been described. The microsomal membranes were resuspended in ⁵⁰ mm Tris-HCl (pH 8) containing ¹ mM mercaptoethanol (buffer 1). To this suspension (protein concentration of approximately ⁸ mg/ml) Emulphogene BC 720 was added, so as to obtain a detergent to protein ratio of 2 (w/w). This emulsion was mixed for 15 min at 30°C, then centrifuged at 100,000g for ¹ h.

¹ Part III in the series "Phospholipid-dependence of plant UDP-glucose sterol β -D-glucosyl transferase" (5, 29).

² Abbreviations: SG, steryl glucosides; ASG, acyl steryl glucosides; UDPG, UDP-glucose; UDPG-SGTase, UDP-glucose sterol β -D-glucosyl transferase; PL, phospholipids; PA, phosphatidic acid; PC, phosphatidylcholine; PE, phosphatidylethanolamine; PG, Phosphatidyl-glycerol; PS, phosphatidylserine.

Ion Exchange Chromatography. A DEAE-cellulose DE-52 (Whatman) column (1 cm i.d.) was prepared and equilibrated in buffer 2 (= buffer 1 containing in addition 20% glycerol $[v/v]$ and 0.2% Emulphogene [w/v]). The supernatant was loaded onto the column (3-4 mg protein/ml gel, flow rate: 0.25 ml/ min), washed with buffer 2 (flow rate: 0.5 ml/min), and then eluted with a discontinuous gradient of KCI (0.050, 0.075, 0.100, 0.150, and 0.400 M KCI in buffer 2; for the profile of this gradient, see Fig. 1). The purification was carried out at 4°C.

UDPG-SGTase Assay. The enzymic preparations (70 μ l of microsomal suspension or Emulphogene supernatant, both diluted 5 times, or 70 μ l of the column fractions) were preincubated for 15 min at 30°C with 20 μ l of (a) an Emulphogene solution (1.8% for microsomes, 1.1% for the supernatant and the DEAE fractions, so as to get the same final detergent concentration of 0.36% w/v); or (b) a sitosterol emulsion in Emulphogene (1.8 or 1.1%) so as to obtain 290 μ M sterol in the final assay mixture except in the kinetic studies where this concentration was varied; or (c) an emulsion in Emulphogene of both sitosterol and different PL at various concentrations, usually 25 μ g/100 μ l final assay (*i.e.* about 300 μ M); we checked that even at the highest sterol concentration used (290 μ m), this substrate is totally solubilized by 0.36% Emulphogene, PL being present or not. Then UDP- $[14C]$ glucose (60,000 cpm in 10 μ l of 50 mm Tris-HCl [pH 8] final concentration in the test usually 0.17 mm except in the kinetic studies) was added and the reaction was run as previously described (5).

The ['4C]glucosylated sterols were then extracted by addition of chloroform:methanol:water so as to get a final ratio of 4:1:1 $(v/v/v)$. The water phase was discarded and the organic phase was washed with methanol:water (1:2, v/v), dried on Na₂SO₄, and transferred into a scintillation vial. The solvent was evaporated and the scintillation cocktail was added (5). It must be recalled that during incubation of the unpurified enzymic preparations, the sterols are not only glucosylated, but part of the SG is acylated into ASG. For this reason we considered the sum of the radioactivities associated with SG and ASG (both extracted

by chloroform-methanol) as a measure of UDPG-SGTase. When separation of ['4C]SG and ['4C]ASG was needed for a measure of steryl glucoside acyltransferase, TLC was performed as described previously (5).

UDPase Assay. This activity was assayed by phosphate determination as described (21).

Phospholipid and Protein Determination. Total lipids were extracted according to Bligh and Dyer (4). All lipids, commercial or extracted, were stored in benzene:ethanol (4:1, v/v) under Ar at -20° C. Mineralization and phosphorus determination were performed as described by Rouser et al. (22). Proteins were assayed according to Schacterle and Pollack (25).

Kinetic Studies.UDPG-SGTase activity was measured with varying concentrations of UDPG in the presence of different fixed concentrations of sterol and vice versa. In both cases the unlabeled UDPG was varied whereas the UDP-[¹⁴C]glucose amount was fixed (60,000 cpm/assay). The UDPG conversion yield was kept below ¹⁰ to 20% and the reaction rate was checked to be constant during the incubation (15 min). The added PL was usually soybean PC (50 nmol).

For the product-inhibition studies the concentration of one substrate was varied in the presence of the second substrate at different fixed (nonsaturating) concentrations and in the absence or in the presence of one product at different fixed concentrations. The product concentrations were of the order of magnitude of the concentrations inducing a 50% inhibition of the enzymatic activity ($I_{50\text{-}UDP}$ approximately 70 μ M and $I_{50\text{-}SG}$ approximately 600μ M). Data were fitted using the iterative nonlinear regression methods of Cleland (7). The programs were written in Fortran and run on a DEC PDP-11 computer.

RESULTS

Solubilization of UDPG-SGTase. Triton X-100 was known to solubilize and activate efficiently plant microsomal UDPG-SGTase (5, 30). Here we used a structurally close detergent which does not absorb at 280 nm: Emulphogene BC 720. At a detergent to protein ratio of 2, 95% of the microsomal UDPG-SGTase was

FIG. 1. Partial purification of UDPG-SGTase and separation from UDPase on ^a DEAE cellulose DE ⁵² column. Typical experiment where ^a ¹⁴ ml sample containing 66 mg protein was chromatographed on a 1×14 cm column as described under "Materials and Methods." Fraction size, 5 ml except for the very first ones. Volumes of the KCI steps: 0.05 m, 45 ml; 0.075 ml, 25 ml; 0.1 m, 25 ml; 0.15 m, 35 ml; 0.4 m, 30 ml.

recovered in a 100,OOOg supernatant (i.e. solubilized) with 65% of the total proteins.

Ion Exchange Chromatography. The 100,OOOg supernatant was fractionated by anion exchange chromatography on a DEAE-cellulose column. The bulk quantity of protein and phospholipid passed straight through the column (Fig. 1). Sterols were similarly eluted (data not shown). A discontinuous KCI gradient was then applied and UDPG-SGTase activity was mainly eluted by 0.15 M KCI whereas UDPase activity was chiefly eluted with 0.05 M KCI and virtually absent from the main UDPG-SGTase peak (fractions 30-38 in Fig. 1). Nor was SG acyl transferase detected in these active fractions (data not shown). The traces of sterol present in these fractions (measured according to Ref. 3), i.e. 0.7 μ g/ml, represent about 1.5 μ M endogenous sterol in the assay mixture (whereas $K_{m\text{-stero}}$ equals approximately 200 μ M).

Purification-Associated UDPG-SGTase Inhibition, and Restoration of Activity by Coarse Lipidic Extract. Table ^I shows the delipidation-associated inhibition of UDPG-SGTase activity, and the stimulation obtained by the simultaneous addition of

Table I. Partial Purification and Delipidation of UDPG-SGTase, after Chromatography on ^a DEAE-Cellulose DE 52 Column

Results of a typical experiment (see also Fig. 1). The experimental error is $\pm 10\%$.

^a UDPG-SGTase activity was measured in the absence or in the presence of lipids: sterols (12 μ g sitosterol, *i.e.* 290 μ *M*, final concentration in the assay) + a lipidic extract of maize coleoptiles microsomes (extraction according to Bligh and Dyer (4); the added extract represents approximately 10 μ g sterols and 50 μ g PL). \blacksquare In this experiment, the purification factor was 7.5. According to the experiments, it varied between 6 and 10. The total activity in fractions 30-38 represented 45% of the total activity loaded onto the column (in cpm). This yield varied between 40 and 70. ^c In fact sterols by themselves partially revealed the enzymic activity and the subsequent addition of the lipidic extract brought about a further stimulation by a factor 2.

Table II. Reactivation Factor of Purified UDPG-SGTase, by Different Phospholipids

The control activity $(= 1)$ was measured in the absence of PL. All the assays (100 μ l) contained: active fraction from DEAE-cellulose (approximately 10 μ g prot), sitosterol (290 μ M), and Emulphogene (0.36%).

 25μ g of these PL were present in the assay (approximately 300 μ M). ^bIn experiments 1 and 2, PC was PC di C_{18:2}; PE and PI were from soybean; PS from bovine brain and PG and PA, from egg yolk PC (via phospholipase D-catalyzed hydrolysis or transesterification).

sterols and a lipidic extract of microsomes. In fact sterols by themselves, as substrates, partially revealed the activity; the sterol amount used was shown to produce the maximal effect; the subsequent addition of the lipidic extract brought about a further stimulation by a factor 2.

Reactivation by Different Pure Phospholipids. We compared the reactivating power of different PL. Table II shows the results of two typical experiments (1 and 2). Whereas the general reactivation amplitude varies from one experiment to the other (more than six were performed), the reproducible feature is the much stronger activation obtained with the negatively charged PL (PA, PG, PI, and PS): they are 2 to 4 times more efficient than the zwitterionic PL (PE and PC).

The fatty acid composition of the natural PL described in experiments ¹ and 2, Table II, is not known. Therefore, we compared the effects of synthetic zwitterionic and negatively charged PL with the same fatty acid composition (Table II, experiment 3). Whatever is this composition, PG is more efficient than PC in stimulating the enzyme.

The importance of the charge of the polar head of PL in the activation of UDPG-SGTase was further confirmed by a comparison of 5 different fatty acids (C_{160} to $C_{18:3}$) with the corresponding methyl esters, for their ability to stimulate purified UDPG-SGTase. The free fatty acids were as efficient as PA in activating the enzyme, whereas the methylesters were one-half as efficient (A Ury, and P Bouvier-Navé, unpublished results).

Kinetic Studies. In order to determine the mechanism for UDPG-SGTase-catalyzed sterol glucosylation and the kinetic parameters of this reaction, we performed kinetic studies with the purified enzyme preparation in the presence of Emulphogene and soybean PC. When UDPG concentration was varied in the presence of different fixed concentrations of sterol (Fig. 2) and vice versa, intersecting patterns were obtained in Lineweaver-Burk plots.

These results clearly indicate that a sequential rather than ping-pong mechanism applies to UDPG-SGTase-catalyzed reaction (27). Thus, they confirm our preliminary data obtained with the acetone powder of maize coleoptiles microsomes (29). The double reciprocal plots and the secondary plots $(1/V_{\text{max}})$ apparent = $f(1/[S])$ allowed the determination of K_m s and V_{max} (mean of two experiments, with PC as the activator): $K_{m\text{-}UDPG}$ = 40 μ M \pm 10%, $K_{m\text{-stero}} = 200 \mu$ M \pm 30%, and $V_{\text{max}} = 160 \text{ pmol}$ / mg prot/s \pm 25%.

Additional insight into the reaction mechanism of UDPG-SGTase was obtained by a study of the inhibition by the reaction products: UDP and SG. Because these products are the substrates of two microsomal enzymes, UDPase (21) and SG acyl transferase (30), these two interfering enzymic activities had to be removed to allow such a study. This was achieved by purification on DEAE cellulose (see Fig. ¹ and the corresponding text). The type of inhibition produced by SG or UDP towards UDPG or sterol was examined both by Lineweaver-Burk plots $(e.g., Fig. 3)$ and by a computer-based statistical analysis of the data, using Cleland's program (7). Whereas both methods failed to determine whether the inhibition by UDP towards UDPG is competitive or noncompetitive, although at least six different experiments were carried out, both methods were in accordance with the exclusion of a competitive type of inhibition for the couples SG/ sterol, SG/UDPG, and UDP/sterol.

Thus, the product-inhibition pattern excludes the possibility of a random bi-bi mechanism for the reaction of sterol glucosylation (27). If UDP is ^a competitive inhibitor towards UDPG, the sterol glucosylation mechanism is simply ordered, UDPG being the first bound substrate and UDP the last released product. If, however, UDP is ^a noncompetitive inhibitor towards UDPG, then the product inhibition pattern is in agreement with a monoiso-ordered mechanism in which a conformational change of the

FIG. 2. Double reciprocal plots of initial velocities with UDPG as the variable substrate and sterol as the changing fixed substrate. \blacksquare [sterol] = 72 μ M (S₁), O [sterol] = 145 μ M (S₂), Δ [sterol] = 290 μ M (S₃). *Inset*: intercept replot.

FIG. 3. Double reciprocal plots of initial velocities with UDPG as the variable substrate, sterol at two fixed concentrations in the absence or in the presence of SG as inhibitor. Noncompetitive pattern. \circ [sterol] = 97 μ M, \bullet [sterol] = 97 μ M and [SG] = 500 μ M, Δ [sterol] = 193 μ M, \blacktriangle [sterol] = 193 μ M and [SG] = 500 μ M.

enzyme is assumed during the course of the reaction (27). In this case the binding order of the substrates (and the release order for the products) cannot be predicted.

Effects of Phospholipids on the Kinetic Parameters of the Reaction. All the kinetic studies described above were carried out in the presence of Emulphogene and soybean PC. In order to evaluate the influence of PL on the K_m s and V_{max} of the reaction, we measured the $K_{m\text{-sterol}}$ (in the presence of 60 μ M UDPG) and $K_{m\text{-}UDPG}$ (in the presence of 290 μ M sterol) of the DEAE-cellulosepurified enzyme (a) in the absence of any PL, or (b) in the presence of PC, or PG (25 μ g). The results show that $K_{m\text{-sterol}}$ is strongly lowered (from 800 to 160 or 140 μ M, respectively) by either PC or PG whereas $K_{m\text{-} \text{UDPG}}$ is only slightly lowered (1.3 times). The apparent V_{max} is also varied by addition of PL, and the observed increase depends on the added PL: when PC increased V_{max} by a factor 1.9, PG increased it by a factor 3.8.

DISCUSSION

Anion exchange chromatography had already been described for partial purification of UDPG-SGTase solubilized from other plant sources (30). Thus, Forsee et al. (10) purified UDPG-SGTase from developing cotton fibers and seeds on a DEAEcellulose and observed a 3-fold increase of the partially purified

enzyme activity, following the addition of purified cotton lecithins. Our results are in agreement with their findings.

We took advantage of the purification-associated delipidation (Table I) of the maize coleoptiles microsomal UDPG-SGTase to confirm its phospholipid dependence (5, 29) and to look for a possible selectivity in this dependence; indeed, UDPG-SGTase exhibited a clear preference for negatively charged PL (Table II). Its behavior is thereby similar to the one found for potato microsomal NADH-Cyt c reductase (20).

The chromatography on DEAE-cellulose allowed a good removal of UDPase and SG acyl transferase from the UDPG-SGTase active fractions. Hence, a complete kinetic study was rendered possible. The data obtained in the absence of products confirm that UDPG-SGTase-catalyzed reaction follows a sequential mechanism (29).

The product inhibition study indicates an ordered addition of the substrates (and removal of the products) with or without an isomerisation of the enzyme. These results are in agreement with the kinetic mechanisms established for two soluble glucosyl transferases (18, 26).

Because UDPG-SGTase is a membrane-bound enzyme, its purification had to be carried out in the presence of detergent. Moreover, one of the substrates (sterol) and one of the products (SG) are insoluble in water. Subsequently the kinetic studies were also carried out in the presence of detergent, *i.e.* with an isotropic system. However, the kinetic parameters of the reaction as well as the enzymic selectivity towards PL might be modified in this micellar system, as compared with ^a membrane system. We intend to perform these measurements in a reconstituted system consisting of the enzyme embedded into liposomes of controlled sterol and PL composition. In spite, or because, of these difficulties inherent to the hydrophobic character of the protein, the knowledge of the kinetic mechanism of the reaction-and of its overall molecular mechanism, including the PL role—is important. Up to nowjust a few plant membrane-bound enzymes have been studied as for their kinetic mechanism (1, 2). No PL dependence was described in these cases.

In order to approach the molecular basis of the activation by PL, we studied their effect on the kinetic parameters of the reaction. We showed that PL affect both the K_m s and the V_{max} of the UDPG-SGTase-catalyzed reaction. $K_{m\text{-sterol}}$ is much more strongly modified than $K_{m\text{-} \text{UDPG}}$ and is 5 times lower with either PC or PG than without. In addition, these PL affect the V_{max} with PG increasing it much more than PC.

We cannot exclude the possibility that PL might be interacting with unidentified inhibitory molecules in the preparations rather than directly with the enzyme. Such inhibitors would be present in the native membrane but kept aside from the enzyme by PL (even after solubilization). They should then co-migrate with the enzyme during the purification step, and be brought into contact with the enzyme, via the purification-associated delipidation. Exogenous PL added to the purified enzyme would then again discard the inhibitory molecules. Although we have never heard of such a situation for any membrane-bound enzyme described in the literature, this possibility remains.

On the other hand, the possibility of a PL role restricted to the 'solubilization' of hydrophobic enzymatic substrates has been reviewed, but in this case usually there is no selectivity towards any PL and even detergent can substitute for PL (23). It must be pointed out here that detergent is present in our assay but is unable in the presence of sterols, to reveal the full activity of the delipidated enzyme (Table II; also Refs. 5, 29). However, this possibility of a mere solubilizing role for PL was evoked for UDPG-SGTase (17). In order to test this hypothesis we incubated our DEAE-cellulose-purified enzymic preparation in the presence of labeled UDPG and different sterol concentrations (from 290 μ M down to 2.9 μ M) in the absence or the presence of PG at different concentrations. The results (data not shown) indicate that whatever the sterol concentration is, UDPG-SGTase activity is stimulated by PG. Moreover, the stimulation factor increases when the sterol concentration decreases. Therefore, we believe that negatively charged PL play much more than a simple role of solubilizing the hydrophobic substrate of UDPG-SGTase. In this respect the possibility of a lowering of the activation energy of the reaction should be considered; this has been clearly shown in the case of animal liver UDP-glucuronyl transferase (13, 19).

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