

PURIFICATION AND CHARACTERIZATION OF UDP-GLUCOSE:SINAPIC ACID GLUCOSYLTRANSFERASE FROM BRASSICA NAPUS SEEDLINGSShawn X. Wang¹, Thomas Vogt² and Brian Ellis¹¹ Department of Plant Science, University of British Columbia, Vancouver, BC, V6T 1Z4 CANADA² Institute of Pflanzenbiochemie, Halle, Germany**ABSTRACT**

UDP-glucose:sinapic acid glucosyltransferase (SGT) is one of the key enzymes involved in biosynthesis of sinapine in *Brassica napus* and other cruciferous species. SGT has been purified 134-fold from 65-h-old seedlings of *B. napus* cv. Westar and found to have a native Mr of 42,500 and a pI of 5. At its optimum pH of 6.0, SGT showed a Km for UDPG of 0.24 mM and for sinapic acid of 0.16 mM. The enzymatic mechanism fits the random bi-bi mode. SGT activity could be detected in all growth stages of *B. napus* plants, but was most active in juvenile tissues and developing seeds.

INTRODUCTION

Sinapine, a phenolic ester of sinapic acid and choline, makes up about 2% of the air-dried oil-free canola/rapeseed meal. Elimination of sinapine from the seeds would improve the flavor, palatability and nutritional properties of canola meal and thereby enlarge markets for canola. Sinapine content in seed may vary from species to species, and from cultivar to cultivar within *Brassica* (Mueller et al. 1978; Wang, 1992) but variation is quite limited (17.3 - 21.6 mg/g meal). An alternative route to low sinapine germplasm is through genetic engineering. The biosynthesis of sinapine is a secondary metabolic process during seed development. Enzymatic synthesis studies in *Raphanus* indicated that sinapic acid could not be directly used to synthesize sinapine. Sinapic acid had to first be activated by the formation of the glucose ester of sinapic acid (1-O-sinapoyl- β -D-glucose, SinG) from UDPG and sinapic acid. This reaction is catalyzed by UDPG:sinapoyl glucosyltransferase (Strack, 1980; Nurmam and Strack, 1981). The SinG provides the acyl donor for the synthesis of sinapine, which is catalyzed by SinG:choline sinapoyltransferase (SCT). Our ultimate objective is to use antisense technology to create a new genotype of *B. napus* with a greatly reduced content of sinapine in the seeds. As the first step towards gaining access to the SGT gene, we are purifying SGT and investigating SGT expression during plant development.

MATERIALS AND METHODS

B. napus cv. Westar was obtained from Agriculture Canada, Saskatoon Research Station. Sinapic acid was purchased from Aldrich Chemicals. All of other chemicals was from Sigma. The 65-h-old seedlings were frozen with liquid nitrogen and then ground with a coffee mill. The fine powder was extracted with chilled 50 mM Tris buffer, pH 7, including 5% glycerol, 4% PPVP and 10 mM 2-mercaptoethanol (2-ME). Desalted protein was used for a series of chromatography (see Table 1). All buffers contain 5% glycerol and 10 mM 2-ME or 2 mM DTT. Westar plants grown in the greenhouse were separated into different tissue types and analyzed at all growth stages. Tissue samples were extracted with Tris buffer. After removing tissue residue by centrifugation, the supernatant was used for SGT activity assay. The SGT activity was assayed by HPLC on a 0.46 x 250 cm Hypersil RP MOS 5 μ HPLC column (Alltech) and the eluent absorbance was measured at 350 nm.

RESULTS AND DISCUSSION**Purification**

SGT from 65-h-old *B. napus* seedlings was purified 134-fold with an overall yield of 0.5% according to the 9-step purification procedure listed in Table 1. The purified SGT was not homogeneous at this point. The protein profile on a non-SDS 10% PAGE gel showed one major and two minor protein bands after silver staining.

Table 1. Purification of SGT from 65-h-old seedlings of *B. napus* cv. Westar

| Purification step | Total Activity (pKat) | Specific Activity (pKat/mg protein) | Total protein (mg) | Enriched fold |
|------------------------------|-----------------------|-------------------------------------|--------------------|---------------|
| Crude extract | 42743 | 19.4 | 2200 | 1.0 |
| Protamine sulfate | 41353 | 28.2 | 1465 | 1.5 |
| 60% Ammonium sulfate | 15380 | 59.8 | 257 | 3.1 |
| Prep-Mono Q (anion exchange) | 11571 | 100.6 | 115 | 5.2 |
| Hydroxyapatite | 8762 | 190.5 | 46 | 9.8 |
| EAI-SinA affinity | 6482 | 308.7 | 21 | 15.9 |
| Mono Q (anion exchange) | 5714 | 317.4 | 18 | 16.3 |
| Mono P (Chromatofocusing) | 3651 | 688.9 | 5.3 | 35.5 |
| Superose 12 (gel filtration) | 1373 | 1961.4 | 0.7 | 101.0 |
| Reverse immunofluorescence | 235 | 2611.1 | 0.09 | 134.4 |

Physical and chemical properties

The general characteristics of partially purified *Brassica* SGT are summarized in Table 2. This enzyme can catalyze the reaction for both directions: SinA + UDPG \leftrightarrow SinG + UDP. Most of *Brassica* SGT characteristics are similar to the enzyme isolated from *R. sativus*. However, the kinetic properties of SGT from *B. napus* suggests that its catalytic mechanism fits the random bi-bi model, whereas the enzyme from *R. sativus* appeared to use an ordered bi-bi mechanism (Moek and Strack, 1993).

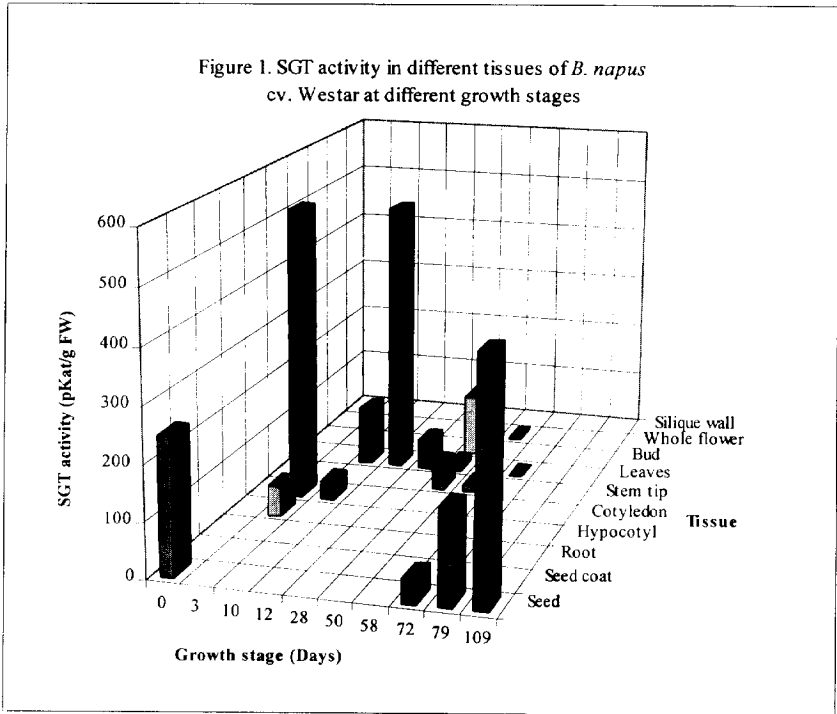
Table 2. Characteristics of SGT from 65-h-old seedlings of *B. napus* cv. Westar

| Item | Properties |
|--------------------------------|--|
| MW | 42.5 KDa (Native) |
| Isoform | Not detected |
| pI | pH 5 |
| Optimum pH | pH 6 (100 mM MES), active at pH 5 - 8 and stable at pH 6 - 7. |
| Optimum temperature | 32 °C, active at 4 - 42 °C and stable at < -20 °C. |
| Co-factor | Not detected |
| Divalent ion | Not required, sensitive to Zn ⁺⁺ , Cu ⁺⁺ , Hg ⁺⁺ , Fe ⁺⁺ , Co ⁺⁺ , inhibition by Zn ⁺⁺ , Co ⁺⁺ and Fe ⁺⁺ was reversible with EDTA, but not Cu ⁺⁺ and Hg ⁺⁺ |
| Reducing reagent | DTT, 2-ME required for stability, but not ascorbic acid. |
| Inhibitor | PHMB showed inhibition, but not NEM, IAA |
| Analogue inhibition | UDP, TDP, UDP-mannose showed strong inhibition. |
| Relative substrate specificity | Sinapic acid 100, ferulic acid 77, 5-OH-ferulic acid 39, cinnamic acid 24, p-cumaric acid 21, caffeic acid 14, syringic acid 10 |
| Reaction mechanism | Random bi-bi, K _[UDPG] 0.24 mM, K _[sinapic acid] 0.16 mM, V _{max} 10.6 pKat |
| Reversibility | UDP, TDP can be used to form UDPG and TDPG. |
| Subcellular location | Cytosol |

Developmental expression of SGT

The developmental expression of SGT of *B. napus* cv. Westar is summarized in Figure 1. SGT specific activity in whole seedling tissue reached its peak between the 2nd and 3rd day, then decreased to 80%. This indicates that the SGT gene is highly expressed in the early stages of seed germination. The physiological function of SGT at this stage is very clear. Highly activated or newly synthesized SGT can rapidly convert sinapic acid released from sinapine to sinapoylglucose. When true leaves developed, the SGT level in the cotyledons declined dramatically. Stem tissue did not show SGT activity except at the apex. In mature plants, most of the SGT activity was found in the leaves; younger leaves showed higher SGT activity than older leaves but no activity in the petiole tissue. When *B. napus* plants went into reproductive stage, the majority of SGT activity was found in the floral bud and stem apex. During the seed

development stage, seed tissue was the only part of the plant which showed detectable SGT activity. SGT specific activity (pKat per mg of protein) reached its peak when the seed was green (after 26 days of flowering). Total SGT activity (pKat per g of fresh seed) continued to increase, but the specific activity gradually declined after the green seed stage, probably due to seed dehydration and accumulation of storage proteins.



ACKNOWLEDGMENTS

The authors thank the Natural Science and Engineering Research Council of Canada for the financial support.

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