AGLYCONE FRAGMENTATION ACCOMPANIES & GLUCOSIDASE CATALYZED HYDROLYSIS OF SALICORTIN, A NATURALLY-OCCURRING PHENOL GLYCOSIDE

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Summary: Salicortin (1a), upon enzyme-mediated glycosidic hydrolysis, undergoes an unusual and rapid fragmentation-recombination reaction leading to 2 via postulated ortho-quinone methide (3) and enol (4) intermediates.

Phenol glycosides are widely distributed plant secondary metabolites that are generally thought to play an important role in regulating herbivory by insects and mammals.¹ In vitro, glucosidase treatment of phenol glycosides provides a mild and efficient method for releasing intact aglycones. In contrast, we report here that enzymatic hydrolysis of salicortin (1a), ² a phenol glycoside that is present³ in many members of the Salicaceae family and that is toxic to some herbivores, ^{4,5} releases glucose³ and leads to a unique fragmentation-recombination reaction of the aglycone that probably involves a transient orthoquinone methide.

Product studies were carried out with substrate levels of β -glucosidase (1 mg mL⁻¹ enzyme,⁶ 5mM 1a, pH 5 acetate buffer, 37 °C). Under these conditions 1a was consumed, carbon dioxide was produced (75% yield by respirometry), and one major product (2)⁷ was formed (53% yield by GC) (Scheme 1). No aglycone (1b) was detected.



Formation of 2 can be rationalized (Scheme 2) by initial hydrolysis of 1a to form 1b. The aglycone (1b) then fragments to ortho-quinone methide 3, carbon dioxide, and an enol (4), which subsequently combines with 3 to form 2. The recombination can be envisioned either as a Michael addition⁸ or as a heteronuclear Diels-Alder reaction⁹ that forms a hemiketal which subsequently opens to give 2.

Trace amounts of o-hydroxybenzyl alcohol, 6-hydroxycyclohex-2-ene-1-one, and catechol were also identified as reaction products by comparison of the gc-ms of the reaction mixture with authentic compounds. These products may arise from solvent capture, tautomerization, or air oxidation of the fragmentation products, respectively.

β-Glucosidase-catalyzed decomposition of 1a is accompanied by a slow time-dependent irreversible inhibition of the enzyme. In a typical reaction at pH 6.2, 25 °C, and 4.4 mM 1a, the half-life of β -glucosidase activity was 70 min. Enzyme activity could not be restored by repeated ultrafiltration with fresh acetate buffer. The inactivation process, which is still under investigation, may be due to reaction of the β -glucosidase active site with the electrophilic 3.

The reaction described here is significant for several reasons: 1) it is an unusual enzyme-catalyzed fragmentation-recombination; 2) it represents a new approach to the development of inhibitors of glucosidases for the rapeutic purposes and as probes for the glucosidase active site; 10 and 3) it may explain the reported toxicity of certain phenol glycosides¹ and in particular the sensitivity of insects having high endogenous levels of β-glucosidases to dietary 1a.⁴

Acknowledgements

We thank the National Science Foundation for support of this research (Grant No. BSR 8416461 and BBS-8614587). The mass spectral determination was performed by the Midwest Center for Mass Spectroscopy, an NSF Regional Instrumentation Facility (Grant No. CHE-8620177). Finally, we thank Dr. Larry Byers for his critical review of an earlier version of this manuscript.

References and Notes

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 (d) ICN almond β-glucosidase, 22 units mg⁻¹ (1 unit = 1 μmole salicin hydrolyzed min⁻¹ at 37⁰C).
 (f) ICN almond β-glucosidase, 22 units mg⁻¹ (1 unit = 1 μmole salicin hydrolyzed min⁻¹ at 37⁰C).
 (f) 2: High resolution mass spectra: found 218.0941; C_{1,3}H_{1,0}O₃ requires: 218.0943; Infrared (CHCl₃): 1690 cm⁻¹ (conjugated ketone); ¹H-NMR (CDCl₂; 90 MHz): δ = 8.5 ppm (1H, broad s, exchangeable with D₂O), 6.7-7.3 (5H, m), 6.2 (1H, ddd, J = 10.0, 2.45, 1.46 Hz), 4.85 (1H, broad s, exchangeable with D₂O), 3.05 (2H, s), 2.45-2.75 (2H, m) and 1.7-2.45 (2H, m). The hydrogen at δ = 6.2 was found to be coupled with hydrogens at δ = 7 and 2.5 by double irradiation experiments. ¹³C-NMR (CDCl₂; 22.5 MHz): δ = 156(s), 151.7(d), 131.8(d), 129.1(d), 125.8(d), 121.9(s), 120.2(d), 117.7(d), 77.5(s), 39.8(t), 33.1(t) and 25.0(t).
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