



## The Chemotaxonomic Significance of Root Anthraquinones and Pre-anthraquinones in the Genus *Lomatophyllum* (Asphodelaceae)

BEN-ERIK VAN WYK,\* ABIY YENESEW† and ERMIA S DAGNET

\*Department of Botany, Rand Afrikaans University, P.O. Box 524, Auckland Park, Johannesburg, 2006, South Africa;

†Department of Chemistry, Addis Ababa University, P.O. Box 1176, Addis Ababa, Ethiopia.

**Key Word Index**—*Lomatophyllum*; Alocoideae; Asphodelaceae; roots; anthraquinones; pre-anthraquinones; chemotaxonomy.

**Abstract**—The roots of seven species of *Lomatophyllum* were analysed by TLC and HPLC for the presence of nine different anthraquinones and pre-anthraquinones which are all known to be characteristic constituents of the subterranean metabolism of *Aloe*. Chrysophanol and asphodelin were detected in all the samples analysed. In addition, the 1-methyl-8-hydroxyanthraquinones aloesaponarin I, aloesaponarin II and laccaic acid D-methyl ester, together with the corresponding pre-anthraquinones were detected in all seven species. The results support the inclusion of *Lomatophyllum* in *Aloe*.

### Introduction

*Lomatophyllum* Willd. comprises 12 species confined to Madagascar, Mauritius and the Aldabra Islands (Smith and van Wyk, 1991). The genus is closely related to the genus *Aloe* and the only known diagnostic character is that the fruits are fleshy at maturity in *Lomatophyllum* while they are capsular in *Aloe*. We here present chemotaxonomic evidence that the close similarity between the two genera is more than just superficial and that *Lomatophyllum* may be no more than a natural section of *Aloe*.

### Materials and Methods

**Plant materials.** Nine root samples from seven species of *Lomatophyllum* were collected from the Johannesburg Botanic Garden (JBG) and from the gardens of the National Botanical Institute, Pretoria (NBI), South Africa. Voucher numbers are indicated in Table 1.

**Procedures.** Freshly harvested roots were air-dried and powdered. Approximately 1 g of each sample was extracted in acetone and prepared for TLC and HPLC analyses as described previously (van Wyk *et al.* 1994). Compounds were directly compared with authentic reference samples (the same as in Dagne *et al.* 1994), using TLC spot characteristics and  $R_f$  values (UV light; 254 and 366 nm), as well as HPLC ( $R_f$  values and UV/VIS spectra). Details of the methods used are given elsewhere (van Wyk *et al.* 1994).

### Results and Discussion

The 1,8-dihydroxyanthraquinones chrysophanol (1), asphodelin (2) and the pre-anthraquinone aloechryson (3), as well as the 1-methyl-8-hydroxyanthraquinones aloesaponarin I (4), aloesaponarin II (5) and laccaic acid D-methyl ester (6), together with the corresponding pre-anthraquinones aloesaponin I (7) and aloesaponin II (8) have been reported to be characteristic constituents of the roots of *Aloe* species (Dagne *et al.*, 1994). The distribution of these lipophilic anthranoid aglycones in seven *Lomatophyllum* species is presented in Table 1, together with a summary of the main chemical patterns in *Aloe* so that the two genera can be directly compared.

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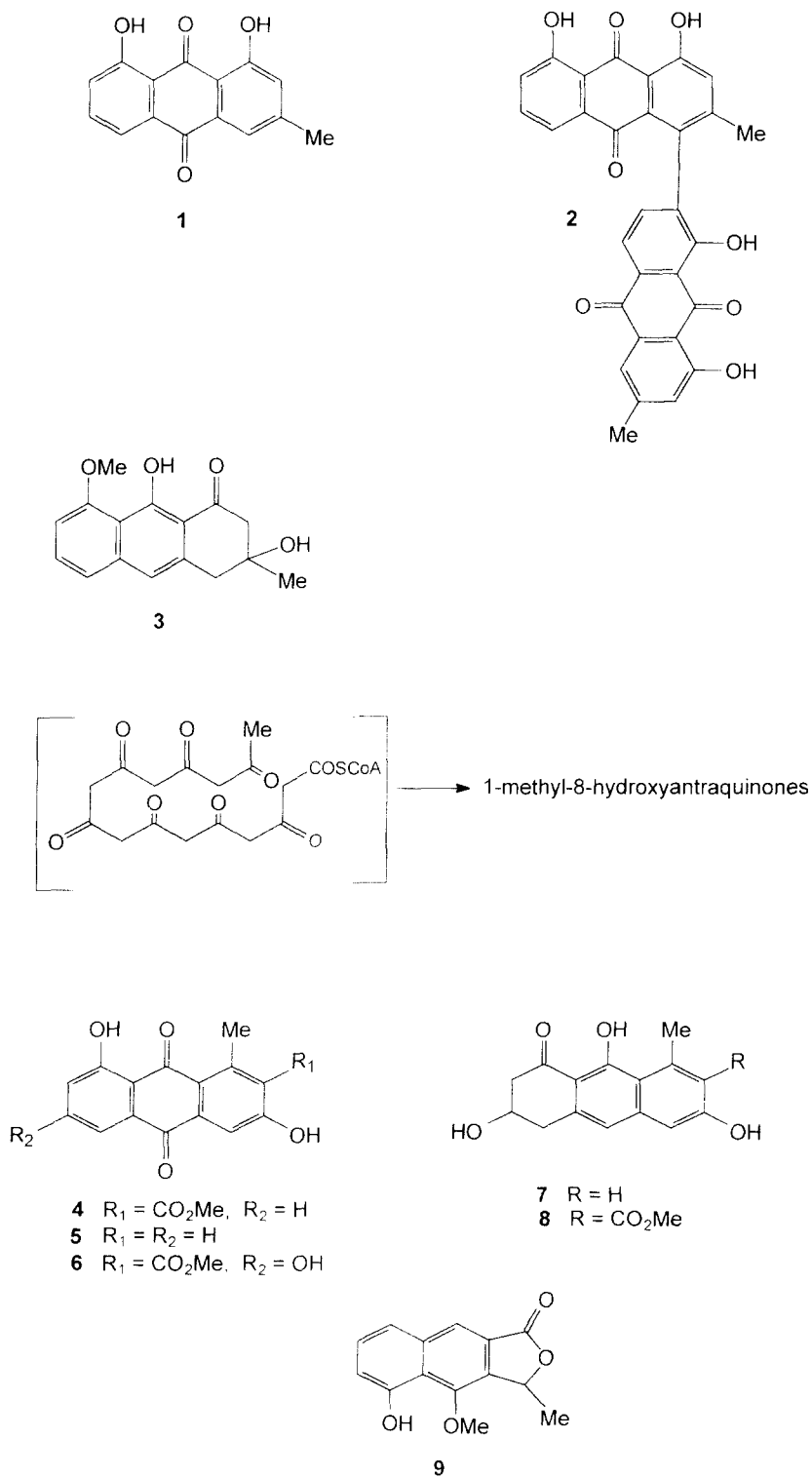


FIG. 1. STRUCTURES OF ANTHRAQUINONES AND PRE-ANTHRAQUINONES IN *LOMATOPHYLLUM*.

TABLE 1. DISTRIBUTION OF LIPOPHILIC ANTHRANOID AGLYCONES (pre-anthraquinones and anthraquinones) IN ROOTS OF SEVEN SPECIES OF THE GENUS *LOMATOPHYLLUM*, COMPARED WITH THE MAJOR PATTERNS IN *ALOE*

Genera and species	Voucher number/ source	Major compounds (see list below)								
		1	2	3	4	5	6	7	8	9
<i>Lomatophyllum aldabrense</i>	NBI 19835	+	+	+	-	+	-	+	+	-
<i>L. antsingyense</i>	ex hort. JBG	+	+	-	-	+	+	+	-	-
<i>L. lomatophylloides</i>	NBI 31156	+	+	+	+	+	+	+	+	-
	NBI 10854	+	+	-	+	+	+	+	+	-
<i>L. macrum</i>	ex hort. JBG	+	+	+	+	+	+	+	+	-
<i>L. occidentale</i>	NBI 31221	+	+	+	+	+	+	+	+	-
<i>L. purpureum</i>	NBI 19135	+	+	+	+	-	+	+	+	-
	NBI 31155	+	+	+	+	+	+	+	+	-
<i>L. cf. purpureum</i>	NBI 17829	+	+	-	+	-	+	+	+	-
Summary of the major patterns in <i>Aloe</i> species (see van Wyk <i>et al.</i> , 1994):										
A. The common pattern, present in 129 out of 172 species sampled		+	+	±	±	±	±	+	+	-
B. Absence of 1-methyl-8-hydroxyanthraquinone pathway, observed in 43 out of 172 species sampled (series <i>Mitriformes</i> , <i>Macrifoliae</i> and other smaller groups, including all tree aloes)		+	+	±	-	-	±	-	-	-
C. Isoeuletherol present (all 16 species of the series <i>Saponariae</i> that were sampled)		+	+	-	±	±	±	+	+	+
D. All nine compounds absent (series <i>Serrulatae</i> )		-	-	-	-	-	-	-	-	-

Compounds: 1, chrysophanol; 2, asphodeline; 3, aloechryson; 4, aloesaponarin I; 5, aloesaponarin II; 6, laccaic acid *o*-methyl ester; 7, aloesaponol I; 8, aloesaponol II; 9, isoeuletherol.

As in *Aloe*, chrysophanol (1) and asphodeline (2) were detected in the roots of all the species of *Lomatophyllum*. These compounds appear to be common in most genera of the Asphodelaceae (Rheede Van Oudtshoorn 1964; Beaumont *et al.* 1985; van Wyk *et al.*, 1994). The pre-anthraquinone aloechryson (3) which was reported to occur sporadically in different groups of *Aloe*, was detected in six of the nine samples.

Interestingly, the 1-methyl-8-hydroxyanthraquinone pathway, which was reported to be characteristic of the genus *Aloe*, have now been observed in all the *Lomatophyllum* species included in this study. Such compounds have not been reported in any other genera of the Asphodelaceae. The occurrence of this pathway in both *Aloe* and *Lomatophyllum* is therefore clear evidence of a close affinity between these genera. Investigations of other genera in the subfamily Alooideae (*Astroloba*, *Chortolirion*, *Gasteria*, *Haworthia* and *Poellnitzia*) will shed more light on the chemotaxonomic significance of the subterranean metabolism at the suprageneric level. On the basis of root anthraquinones and pre-anthraquinones, there are larger discontinuities within *Aloe* than between *Aloe* and *Lomatophyllum* and the generic status of the latter can now be seriously questioned.

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