

Phytochemical, Morphological, and Molecular Evidence for the Occurrence of the Neotropical Liverwort *Plagiochila stricta* in the Canary Islands, New to Macaronesia

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Abstract. *Plant material from the Canary Islands similar to Plagiochila spinulosa (Dicks.) Dumort. has been identified as the Neotropical liverwort P. stricta Lindenb. The secondary metabolite profiles of CDCl₃ extracts of both Neotropical and Macaronesian specimens of P. stricta are distinct from those of P. spinulosa in containing alloocimene and neoalloocimene as the principal monoterpenes rather than β-phellandrene; the two species also show differences in the 9,10-dihydrophenanthrenes and other aromatic compounds present. Phylogenetic analyses of ITS nrDNA sequence variation place P. spinulosa sister to P. stricta. Morphologically, P. stricta from Tenerife is distinguished most easily from P. spinulosa by the scarcely decurrent antical leaf margin.*

There is current interest in the potential of bryophyte studies to provide biogeographical insight into genetic diversity, cryptic speciation, and origins of plant populations, especially on islands within areas such as Macaronesia (Freitas & Brehm 2001), and in the application thereto of phylogenetic analyses of DNA sequence variation or of other DNA-derived data (Shaw 2001). Adequate knowledge of the relevant taxa is essential for such studies and, as in the case of the difficult and numerous genus *Plagiochila*, is not always available.

The bryophyte checklist for the Canary Islands (Dirkse et al. 1993) recorded five *Plagiochila* species on Tenerife: *P. dubia* Lindenb. & Gottsche [now identified as *P. virginica* A. Evans (Heinrichs et al. 2002b)], *P. exigua* (Taylor) Taylor, *P. killarriensis* Pearson [synonymous with *P. bifaria* (Sw.) Lindenb. (Heinrichs et al. 1998b)], *P. punctata* (Taylor) Taylor, and *P. spinulosa* (Dicks.) Dumort. Following our phytochemical studies of *Plagiochila* specimens from Madeira that resulted in the discovery of *Plagiochila retrorsa* Gottsche and of a still unidentified *Plagiochila* with dentate leaves (Rycroft et al. 2001), herbarium specimens of some small plants from the Canary Islands that had been

determined as *P. spinulosa* were analyzed phytochemically. The NMR profiles were different from those of specimens of *P. spinulosa* from the British Isles (Connolly et al. 1999), but had similarities to those of *P. retrorsa* Rycroft et al. 2001) and *P. papillifolia* Steph. (Heinrichs et al. 2002c) in respect of the presence of compounds **1** and **4** (chemical compounds and structures are identified by numerals printed bold in Fig. 1).

Against this background, stands of a *Plagiochila* that superficially resembled *P. spinulosa* (with dentate, sometimes caducous leaves) were observed in three areas of humid laurel forest during the British Bryological Society Spring 2001 Meeting on Tenerife (Blockeel 2002). However, compared to British material, the facies of the plant was different and the odor of the crushed plant in the field was spicier. In addition, the preferred habitat appeared to be excessively shaded compared to that of *P. spinulosa* in the British Isles.

These observations prompted us to investigate the plants further—we were able to capitalize on phytochemical and molecular data that we had already obtained from Neotropical material of *P. stricta* Lindenb., a species whose morphology was

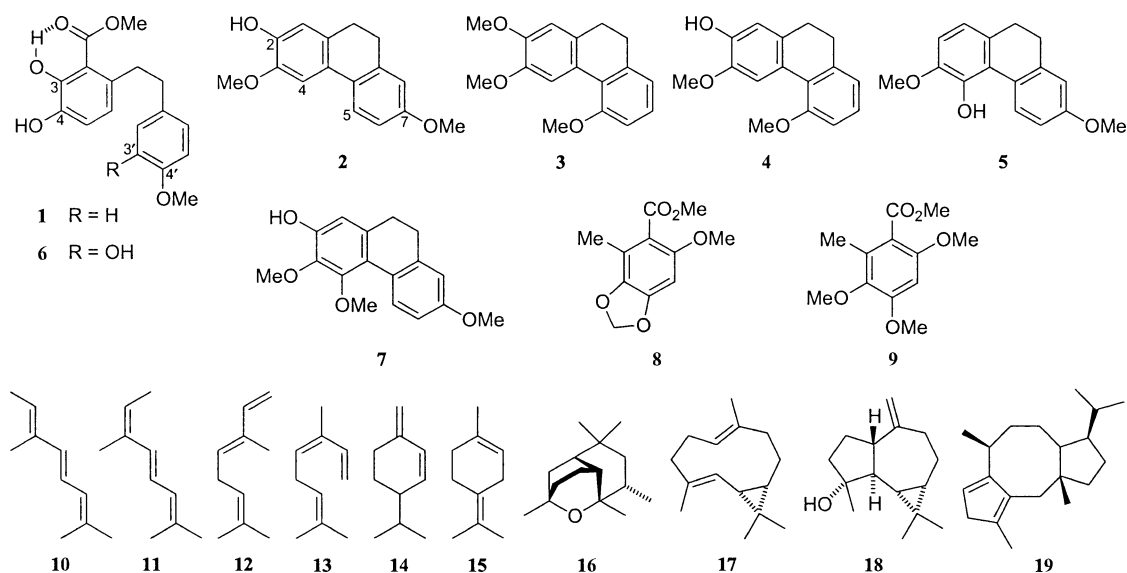


FIGURE 1. Chemical structures, identified by the bold numerals that are used in the text and Table 1.

recently redescribed (Heinrichs et al. 1998a), and to extend our knowledge of the molecular phylogeny of the genus *Plagiochila* (Heinrichs et al. 2002a,b; Renker et al. 2002). Here we report details of work that established the presence of *P. stricta* in the Canary Islands archipelago.

MATERIALS AND METHODS

Plant material.—Details of four of the specimens used are in the caption to Table 1. A fifth, from which DNA was extracted, was *Drehwald 3920*, Tenerife, 24th December 1999 (GOET). Other material used for illustration is mentioned in the caption to Figure 2.

Phytochemistry.—The lipophilic secondary metabolites of *P. stricta* specimens were determined using NMR and GC–MS profiling of deuteriochloroform extracts (Rycroft 1996, 1998). Upper parts of a few shoots were selected and inspected carefully to remove other species. The dry plant material was ground together with a small quantity of broken glass and extracted with CDCl_3 (ca 0.7 ml). Proton NMR spectra of the filtered solutions were measured at 400 MHz, with chemical shifts (δ ppm) reported relative to internal tetramethylsilane. Concentrations were measured by comparing signal integrals with the calibrated residual proton signal of the solvent and converted to weight percentages extracted from the dry plant. The same solutions were used for GC–MS, as described previously (Rycroft & Cole 2001). GC–MS of the extracts after acet-

TABLE 1. Compounds in CDCl_3 extracts of *Plagiochila stricta*. The structure numbers correspond with Fig. 1. The specimens (all in GOET) are i): *Rycroft 01035*, Tenerife, 19 February 2001, 30 mg extracted 27 February 2001; ii): *Rycroft 01071*, Tenerife, 22 February 2001, 35 mg extracted 26 July 2001; iii): *Holz EC-01-478*, Ecuador, May 2001, 23 mg extracted 5 June 2001; iv): *Heinrichs et al. 4401*, Costa Rica, September 1999, 17 mg extracted 16 May 2000. The absolute amounts of compounds extracted from the specimens were determined from proton NMR spectra and converted to % w/w of the dried specimens extracted. The amounts are indicated as follows: “–”: not detected; “tr.”: trace; “+”: 0.04–0.1%; “++”: 0.2–0.9%; “+++”: 1–2%; “++++”: 3–4%; parentheses denote detection by GC–MS only.

Compound	Structure	Specimen number			
		(i)	(ii)	(iii)	(iv)
methyl 4-hydroxy-4'-O-methylunlunurate	1	++++	++++	++++	++
3,7-dimethoxy-9,10-dihydrophenanthren-2-ol	2	+	+	++	+
2,3,5-trimethoxy-9,10-dihydrophenanthrene	3	+	+	+	+
3,5-dimethoxy-9,10-dihydrophenanthren-2-ol	4	++	++	–	++
3,7-dimethoxy-9,10-dihydrophenanthren-4-ol	5	++	++	++	–
methyl 3',4-dihydroxy-4'-O-methylunlunurate	6	(tr.)	(tr.)	++	+
alloocimene	10	+++	++	++	++
neoalloocimene	11	+++	++	+++	++
terpinolene	15	(+)	(+)	(+)	(+)
peculiaroxide	16	++	++	+	++
bicyclogermacrene	17	+++	++	++	+
spathulenol	18	–	++	+	+
fusicocadiene	19	(+)	–	(++)	(++)

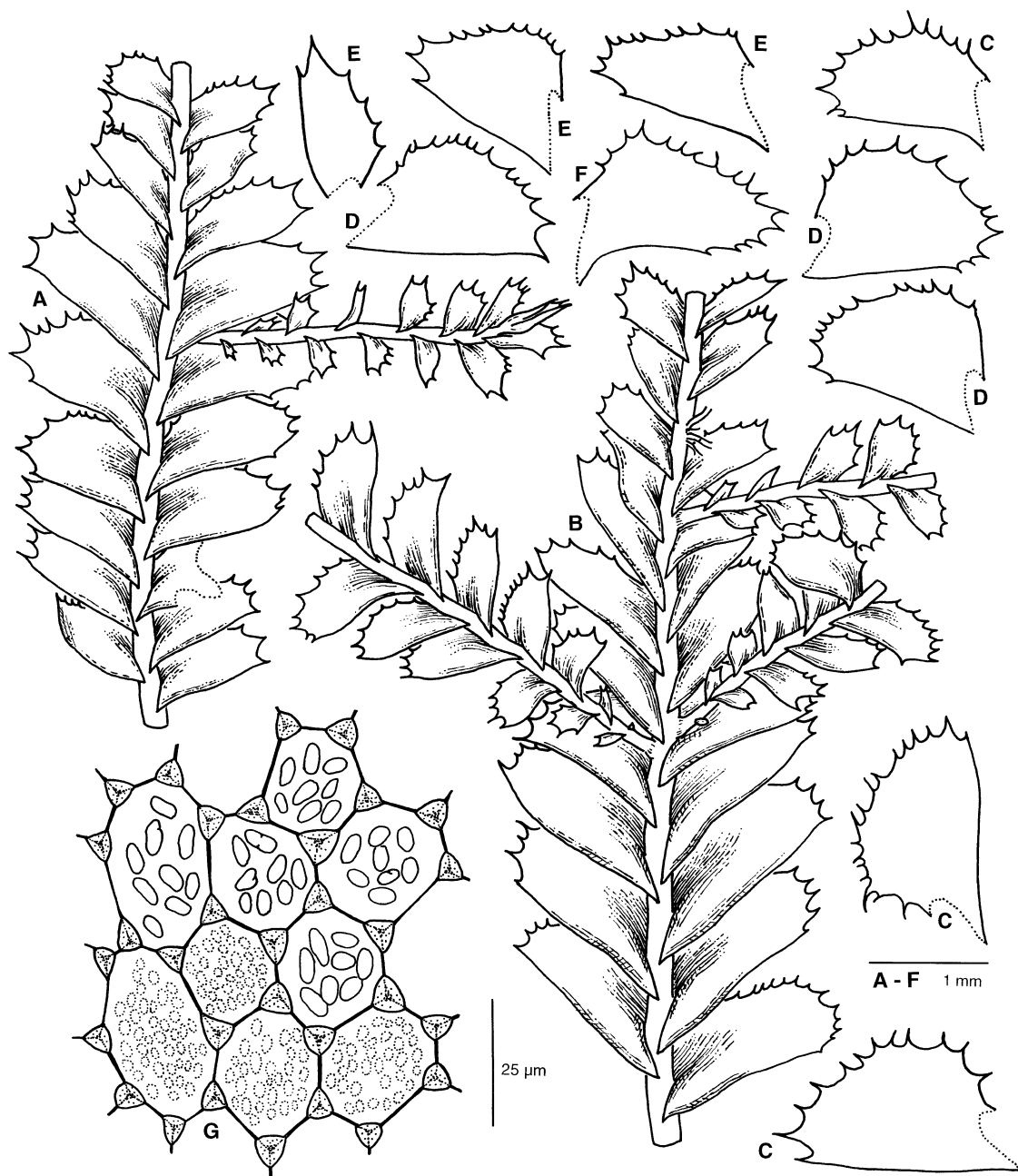


FIGURE 2. *Plagiochila stricta* Lindenb. — A, B. Parts of shoots, dorsal view. — C–F. Leaves. — G. Cells from leaf center, with oil bodies or papillae. (A, D, G from Tenerife, *Rycroft 01071* [GOET]; B, C from Tenerife, *Drehwald 3929* [GOET]; E from Jamaica, *Underwood 1410* [GOET]; F from Ecuador, *Holz EC-01-478* [GOET]).

ylation was also undertaken in order to distinguish compounds with hydroxyl groups that could be acetylated. Compounds were identified by comparison with authentic compounds or with data either in the literature or from our earlier work. There is much confusion in the literature concerning the identification and stereochemistry of allocimene (**10**) and neoallocimene (**11**). Although this situation was highlighted and clarified many years ago (Crowley 1968), the problem continues to manifest itself. In contrast to the information presented in a recent com-

pilation of GC–MS data (Adams 1995), allocimene is in fact (4*E*,6*E*)-2,6-dimethyl-2,4,6-octatriene (Combined Chemical Dictionary 2001), whereas neoallocimene is (4*E*,6*Z*)-2,6-dimethyl-2,4,6-octatriene and has the shorter GC retention time, a consequence of its lower boiling point.

DNA extraction, PCR amplification, cloning, and sequencing.—The procedures leading to the sequences of the nrDNA internal transcribed spacer (encompassing the ITS1-, 5.8S- and ITS2-regions, and designated ITS here-

after) were as described previously (Heinrichs et al. 2002b). GenBank accession numbers of the four *P. stricta* sequences are AJ416646 (766 bases, Heinrichs et al. 4401), AJ416647 (764 bases, Holz EC-01-478), AJ416648 (768 bases, Rycroft 01071), AJ416649 (769 bases, Drehwald 3920). Details of the other eleven sequences used have been published (Heinrichs et al. 2002a).

Phylogenetic analyses.—Phylogenetic trees were inferred by using PAUP* (Swofford 2000) to apply distance (neighbor-joining: NJ), maximum parsimony (MP) and maximum likelihood (ML) criteria to a data set of 802 aligned positions of 15 ITS nrDNA sequences (*Herbertus subdentatus*, four specimens of *P. stricta*, and 10 other species of *Plagiochila*; alignment available from J.H.; initially the alignment resulted in 876 positions, but 74, mainly belonging to inserts of the outgroup, were excluded from the analysis, to avoid producing artificially long branches). Modeltest 3.04 (Posada & Crandall 1998) was used to choose the evolutionary model that fits the data best; this program employs two statistics: the likelihood ratio test (LRT) and the Akaike information criterion (AIC, Akaike 1974). The model selected by the hierarchical LRT was the HKY85 model (Hasegawa et al. 1985), with estimates of the transition/transversion ratio and of the gamma shape parameter for among site variation (G) calculated from the data set. The ML method (applied to the HKY85+G model) was used for phylogenetic analyses.

The confidence of branching was assessed using 100 bootstrap resamplings in ML analysis (using the HKY85+G model, with 10 addition-sequence replicates) and 1,000 bootstrap resamplings in distance (NJ method with the HKY85+G model) and unweighted MP analysis (with 10 addition-sequence replicates) of the data set (Felsenstein 1985).

TreeView (version 1.6.2, Page 1996) was used to modify the tree file of the "best tree" in order to generate user-defined trees. To assess user-defined topologies relative to the "best tree," the sequence data file and the tree files were loaded into PAUP* and tested according to Kishino and Hasegawa (1989); comparisons were based on ML (with the HKY85+G model of evolution selected by Modeltest 3.04) and MP criteria.

RESULTS AND DISCUSSION

Morphological differentiation.—*Plagiochila stricta*, typified on Swartz material from Jamaica, is a widespread, Neotropical, montane species. It was redescribed and illustrated recently (Heinrichs et al. 1998a); several heterotypic synonyms were identified and subsequently a further synonym, *P. elegantula* Herzog, has been reported (Heinrichs et al. 2001). Plants from Tenerife are illustrated in Figure 2. All the material of *P. stricta* observed on Tenerife was sterile and generally less robust than that from the Neotropics.

Material like that now identified as *P. stricta* has been determined previously as *P. spinulosa*. The antical leaf margin of *P. spinulosa* is long-decurrent (Paton 1999) whereas in the Tenerife taxon it is barely or not decurrent. In the British Isles, this character would be sufficient to refer the material to *P. punctata*, but on Tenerife that species is typically very robust, with a facies more akin to that of vigorous *P. spinulosa* in the British Isles. The

tendency of many of the shoots to have leaves that lie more or less flat and spread out, rather than tending to be revolute and retrorse as in *P. spinulosa*, contributes to the distinctive facies of *P. stricta* on Tenerife, where it grew in thin mats close to the substrate rather than in thick cushions. The leaf cuticle papillae of both *P. stricta* and *P. spinulosa* may exhibit different shapes, depending on the proportions of the cell dimensions. Distinctly elongate cells tend to have ellipsoidal to striate papillae whereas round to ellipsoidal papillae are typical of more or less isodiametric cells. Papillae in *P. spinulosa* are usually loosely spaced whereas those in *P. stricta* are often densely spaced. Elongate, triangular to ciliate teeth are sometimes present at the postical leaf base of Neotropical *P. stricta* and can be found in Tenerife material, though infrequently. The ill-defined vitta of *P. stricta* is not well developed in Tenerife material. Although perianths were not observed on Tenerife, those occurring in Neotropical *P. stricta* have a mouth with elongate triangular to long ciliate teeth; *P. spinulosa* has short-teeth on the perianth mouth.

Ecological discrimination.—The preferred habitat of *P. stricta* on Tenerife, in contrast to that of *P. spinulosa* in the British Isles (and, indeed, to that of *P. bifaria*, *P. punctata*, and *P. exigua* on Tenerife), appeared to be deeply shaded and humid, in *laurisilva* localities sometimes also occupied by *Heteroscyphus denticulatus* (Mitt.) Schiffn. and *Trichomanes speciosum* Willd., species with habitat requirements that are different from, and much more demanding than, those of *P. spinulosa*. *Trichomanes speciosum* was described recently as "the most shade tolerant of all plant species so far studied" (Johnson et al. 2000) and the sporophyte generation is restricted to sheltered and extremely humid sites (Ratcliffe et al. 1993). While it is not the intention to suggest that *P. stricta* is equally demanding, it was striking that more open positions where there was luxuriant growth of *P. punctata* (and where *P. spinulosa* might have been expected based on experience in the British Isles) were not occupied by *P. stricta*.

Phytochemical distinction.—Table 1 shows the major aromatic and terpenoid compounds observed in the CDCl₃ extracts of four specimens of *P. stricta*. The profile with respect to the aromatic compounds is not uniform, although the two Tenerife specimens are rather similar. This is evident from the proton NMR spectra of the four extracts in Figure 3, where the methoxyl region is presented so as to ease comparison with the same region of spectra of *P. retrorsa* extracts (Rycroft et al. 2001). One *P. spinulosa* extract is included in Figure 3 and is readily seen to be very different in detail, although the same types of compound are among those pre-

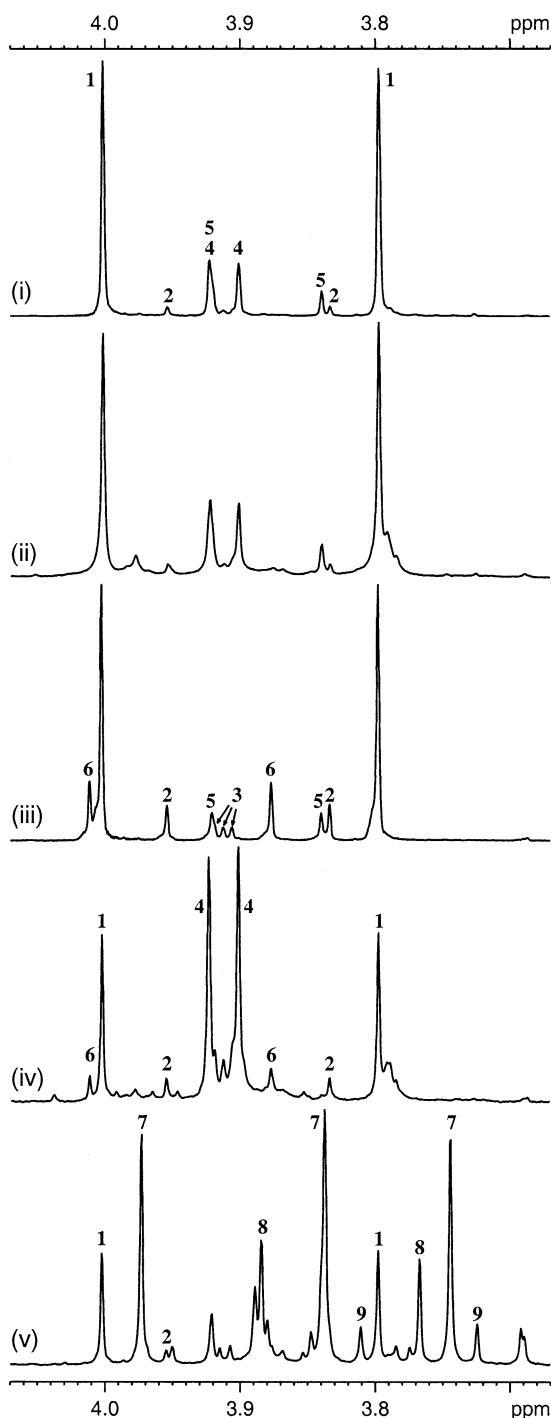


FIGURE 3. The methoxyl region of the 400 MHz ^1H NMR spectra of five CDCl_3 extracts. (i)–(iv) are from the *Plagiochila stricta* specimens in Table 1. (v) is from a *P. spinulosa* specimen for comparison (Rycroft 97137, England). The peak labels refer to structure numbers in Figure 1 and Table 1; for clarity and/or because of overlap, all peaks from a compound may not be labelled.

sent (Connolly et al. 1999). The methyl lunularate derivative **1** and 3,7-dimethoxy-9,10-dihydrophenanthren-2-ol (**2**) are present in all the *P. stricta* extracts and have been reported from *P. spinulosa*, *P. retrorsa* (Rycroft et al. 2001), and *P. papillifolia* (Heinrichs et al. 2002c). 2,3,5-Trimethoxy-9,10-dihydrophenanthrene (**3**), also present in extracts (i)–(iv), is known from *P. retrorsa* and *P. papillifolia* only. 3,5-Dimethoxy-9,10-dihydrophenanthren-2-ol (**4**), reported from *P. retrorsa*, *P. papillifolia*, and *P. bifaria* (Rycroft et al. 1999), but absent from *P. spinulosa*, is present in all but the Ecuadorian specimen (iii). 3,7-Dimethoxy-9,10-dihydrophenanthren-4-ol (**5**), known hitherto from *P. spinulosa* only, is present in all but the Costa Rican specimen (iv).

The NMR spectra of the two Neotropical specimens indicated the presence of an unknown compound, at a level 25% in (iii) and 15% in (iv) of the amount of compound **1**. The unknown compound has one methoxyl group (δ 4.01 p.p.m.) in a similar environment to that of the ester methoxyl of **1** (δ 4.00 p.p.m.), and a second methoxyl (δ 3.88 p.p.m.) deshielded by 0.08 p.p.m. compared to that of the *p*-methoxybenzyl moiety of **1** (δ 3.80 p.p.m.). Corresponding signals in the GC–MS were not observed until these two extracts were acetylated, when a peak of appropriate intensity and with Kováts index 2,854 (compared to 2,619 for the diacetate of compound **1**) gave a mass spectrum with a parent ion at m/z 444 (corresponding to acetylation of three hydroxyl groups in a compound with M_r 318) and a base peak at m/z 137, equivalent to one oxygen atom more than the base peak of compound **1**. It is likely that the unknown compound has structure **6**, a new derivative of **1** where the benzyl moiety responsible for the base peak bears a 3-hydroxyl group as well as a 4-methoxyl. There are ten possible isomers of a hydroxymethoxybenzyl species. Comparison of the chemical shift observed for the methyl ether signal (δ 3.88 p.p.m.) with literature values enables us to reject seven of these immediately, but distinguishing between the 3-hydroxy-4-methoxy-, 4-hydroxy-3-methoxy- and 2-hydroxy-3-methoxy-isomers is less straightforward. The value reported by Crombie and Crombie (1982) for 3,4'-dihydroxy-4,5'-dimethoxybibenzyl (δ 3.88 p.p.m.) fits perfectly and that in 3-hydroxy-3',4'-dimethoxybibenzyl (δ 3.86 p.p.m.) is close (Asakawa et al. 1987). Methoxyl chemical shifts of 4-hydroxy-3-methoxybibenzyls may be as large as δ 3.85 p.p.m., the value reported for gigantol by Tesuka et al. (1991), although smaller values have also been reported for the same compound, e.g., δ 3.82 p.p.m. by Leong et al. (1997); the latter value was reported for 4-hydroxy-3,3'-dimethoxybibenzyl by Anton et al. (1999). Values reported for 2-hydroxy-3-methoxybibenzyls, including δ 3.83 p.p.m. for cirrho-

petalidin (Majumder & Basak 1991) and δ 3.87 p.p.m. for pellepiphyllin (Cullmann et al. 1997), do not allow us to exclude this isomeric possibility. Comparison with chemical shift data in the literature is fraught with difficulties (such as uncertainties over exact measurement conditions, sample concentrations, and chemical shift references, even when these are nominally the same), especially when discrimination between small differences is required: structure **6** remains the most likely candidate but has not been proved.

Several features of the aromatic compound profile distinguish *P. stricta* from *P. spinulosa*. For example, some of the signals labelled in the ^1H NMR spectrum of *P. spinulosa*, Figure 3(v), arise from the major component, 2-hydroxy-3,4,7-trimethoxy-9,10-dihydrophenanthrene (**7**), and two methyl benzoate derivatives (**8** & **9**), compounds that were not detected in *P. stricta*. None of the 9,10-dihydrophenanthrenes in *P. stricta* with an oxygen substituent at position 5 (such as **3** and **4**) is present in *P. spinulosa*.

The monoterpene content is even more distinctive. Analysis soon after collection showed that ca 1% w/w of both alloocimene (**10**) and neoalloocimene (**11**) had been extracted from the plants. Two other ocimene isomers, viz. (*E*)- β -ocimene (**12**) and (*Z*)- β -ocimene (**13**), were detected using GC-MS, but in quantities one and two orders of magnitude less, respectively, than the abundant isomers. These acyclic farnesatrienes are in sharp contrast to the principle monoterpene in *P. spinulosa*— β -phellandrene (**14**), a monocyclic menthadiene; using GC-MS, β -phellandrene was not detected in any of extracts (i)–(iv) of *P. stricta*. This chemical difference is presumably the basis of the field observation that the odor of the crushed plant can be used to distinguish *P. stricta* and *P. spinulosa*. The only other monoterpene in *P. stricta* present in reasonable amount is terpinolene (**15**). There is one previous report of “ocimene” from a liverwort, from *Jungermannia faureana* Beauverd, but the isomeric form was not specified and the structure has been shown both as (*E*)- β -ocimene (Asakawa et al. 1980) and (*Z*)- β -ocimene (Asakawa 1982).

In angiosperms and gymnosperms it is usual that a single enzyme takes the precursor molecule (geranyl pyrophosphate) through to the final monoterpene (Wise & Croteau 1999). Many such monoterpene syntheses have been isolated that yield single or multiple products, although enzymes involved in the biosynthesis of ocimenes are less well known (Bohlmann et al. 2000). It has been suggested that monoterpene biosynthetic reaction mechanisms and enzymes are similar in liverworts and higher plants (Adam & Croteau 1998). If this scenario extends to *Plagiochila*, then our observation of different

monoterpenes in *P. stricta* and *P. spinulosa* suggests the presence of different enzymes and pathways, rather than, for example, loss in *P. stricta* of activity part way along a common pathway.

The remaining compounds in Table 1 are sesquiterpenoids and a diterpene that are commonly found in *Plagiochila* species. The only previous report of chemical constituents of *P. stricta* deals with a specimen from Peru determined as one of the synonyms, *P. acanthoda* Lindenb. & Gottsche (Asakawa & Inoue 1987). No peaks corresponding to any ocimene isomers were among the GC-MS data reported, but monoterpenes are volatile and might easily be lost from voucher specimens. The two most abundant constituents both had parent ions at m/z 256 that were also the base peak and may correspond to 9,10-dihydrophenanthrenes (such as **4** and **5**) with one hydroxyl and two methoxyl substituents.

ITS nrDNA sequence analysis.—The molecular investigation produced a monophyletic lineage (Fig. 4) for *P. bifaria*, *P. punctata*, *P. spinulosa*, and *P. stricta* that is well supported by bootstrap analyses in all methods. The first three of these species were assigned to *P. sect. Arrectae* Carl in an earlier molecular study (Heinrichs et al. 2002b). The analyses also produced a monophyletic lineage for the four *P. stricta* sequences. While the Neotropical and the Macaronesian specimens of *P. stricta* are placed in separate clades with good bootstrap support, the differences are comparable to the variation observed between different stands of other *Plagiochila* species, such as *P. porelloides* (Nees) Lindenb. (Renker et al. 2002) and *P. superba* (Spreng.) Mont. & Nees (Heinrichs et al. 2002a). In view of the lack of substantial phytochemical differences in the morphological and phytochemical data, the monophyletic *P. stricta* lineage is interpreted in favor of a single species.

In order to examine the robustness of branching within the *Arrectae* clade, the topology of the “best tree” (Fig. 4) was compared with user-defined trees (Table 2). KH tests (Kishino & Hasegawa 1989) clearly reject placing *P. bifaria* within the *P. stricta* clade, but do allow *P. punctata* and *P. spinulosa* to be placed there and point to the close relationship of *P. spinulosa* and *P. stricta*. The four *P. stricta* sequences differ in 29 of 802 aligned positions, the Ecuadorian and the Costa Rican specimens differ in 13, and the two Tenerife specimens in eight positions. However, the greater numbers of differences between the four *P. stricta* sequences and the *P. spinulosa* sequence (61 positions) and between the two Tenerife sequences and the *P. spinulosa* sequence (46 positions) justify separation of *P. spinulosa* from *P. stricta*.

Phytogeography.—Until 1998, only two repre-

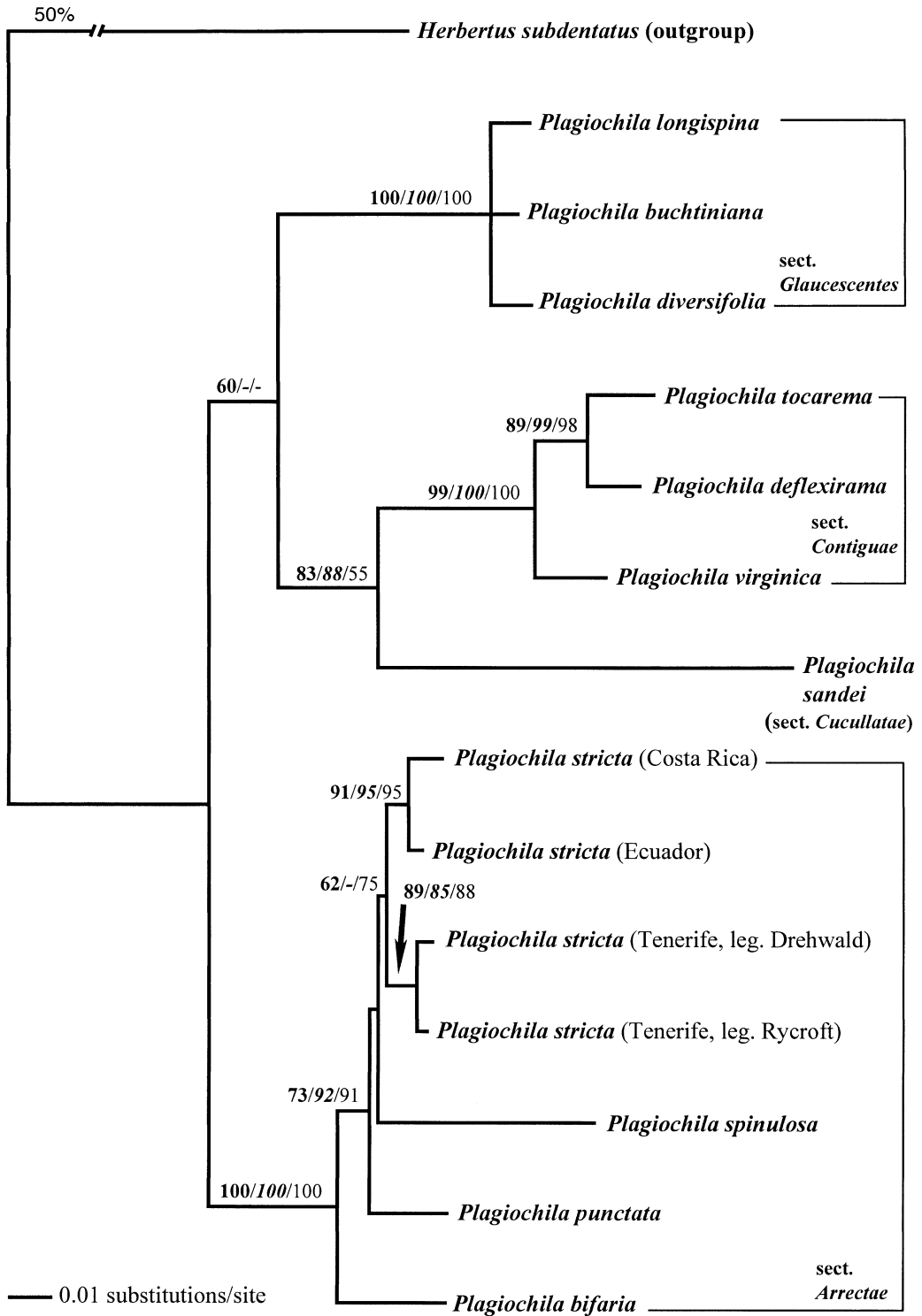


FIGURE 4. Molecular phylogeny of *Plagiochila* species based on ITS1-, 5.8S-, and ITS2-nrDNA sequence comparisons using 802 aligned positions. The rooted tree resulted from a maximum likelihood analysis of 15 sequences (including the outgroup), using the model of Hasegawa et al. (1985) with estimated transition/transversion ratio (Ti/Tv = 1.66) and estimated gamma shape (G = 0.75), HKY85+G, calculated as the best model by Modeltest 3.04 (Posada & Crandall 1998); bootstrap percentage values (> 50%) were determined for maximum likelihood (using HKY85+G; bold), neighbor-joining (using HKY85+G; bold italic) and unweighted maximum parsimony (not bold) methods.

TABLE 2. Comparisons of the maximum likelihood tree in Fig. 4 with user-defined trees by Kishino-Hasegawa tests (one-tailed tests) using maximum likelihood (ML) and maximum parsimony (MP) methods. ^aML using the HKY85+G model after Hasegawa et al. (1985), with estimated transition/transversion ratio (Ti/Tv = 1.66) and estimated gamma shape (G = 0.75); ^b $\Delta(-\ln L)$ = difference in $(-\ln)$ -likelihood between the best tree (Fig. 4) and the user-defined tree; ^cP = probability of getting a more extreme T -value under the null hypothesis of no difference between the two trees (one-tailed test); ^d $\Delta(\text{length})$ = difference in tree length between the best tree (Fig. 4) and the user-defined tree; ^e $(-\ln)$ -likelihood of the best tree (Fig. 4); ^flength of the optimal tree in the maximum parsimony analysis; *indicates that the user-defined tree is significantly worse than the best tree ($p < 0.05$).

Tree topology	ML ^a		MP	
	$\Delta(-\ln L)^b$	P ^c	$\Delta(\text{length})^d$	P ^e
1 best tree (Fig. 4)	(3241.4) ^e	—	(433) ^f	—
2 <i>P. spinulosa</i> as sister to EITHER the two Neotropical OR the two Macaronesian <i>P. stricta</i>	1.0 ± 1.8	0.2783	1	0.3176
3 <i>P. punctata</i> as sister to EITHER the two Neotropical OR the two Macaronesian <i>P. stricta</i>	5.6 ± 4.8	0.1196	2	0.3176
4 <i>P. bifaria</i> as sister to EITHER the two Neotropical OR the two Macaronesian <i>P. stricta</i>	12.9 ± 6.7	0.0283*	8	0.0113*
5 <i>P. bifaria</i> as sister to <i>P. punctata</i>	6.0 ± 4.5	0.0891	5	0.0253*

representatives of the genus *Plagiochila* were recorded from both Macaronesia and the Neotropics, and re-determination has recently removed one of these Neotropical elements from the Macaronesian flora (Heinrichs et al. 2002b). Since 1998, considerable evidence, detailed elsewhere (Heinrichs & Rycroft 2001; Rycroft et al. 2001), has accumulated of close relationships between the *Plagiochila* floras of the Neotropics and Atlantic Europe (including Macaronesia). Determination of material from Tenerife as *P. stricta* has added yet another disjunct Neotropical element to the Macaronesian *Plagiochila* flora.

The observation that there is good bootstrap support for placing the Neotropical and Macaronesian specimens in different clades within *P. stricta* is pertinent to current interest in biogeographic patterns and cryptic speciation in bryophytes (Shaw 2001). While the similarity in morphology and phytochemistry suggest overall that the Neotropical and Macaronesian populations are conspecific (especially in the context of the species concept being applied to the on-going revision of the genus *Plagiochila* in the Neotropics), the available molecular data are also consistent with a hypothesis that there are two cryptic geographical taxa. More sequences are required to decide if the Macaronesian and Neotropical stands are clearly isolated genetically from each other or if the microclades break down when more sequences are included in the analyses.

CONCLUSION

Phylogenetic analyses of ITS nrDNA sequence variation place *P. spinulosa* sister to a clade containing *P. stricta*, and phytochemical data support

separation of the two taxa. We have demonstrated that *P. stricta*, a Neotropical species, occurs on Tenerife in the Canary Islands but the question of whether *P. spinulosa* also occurs on Tenerife (and, more generally, in Macaronesia) requires further investigation.

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