



Botanical Journal of the Linnean Society, 2016. With 6 figures

## Biogeographical, ecological and ploidy variation in related asexual and sexual *Limonium* taxa (*Plumbaginaceae*)

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Received 12 January 2016; revised 26 July 2016; accepted for publication 3 September 2016

Limonium is a widespread genus of halophytes and taxa found on the Atlantic coast include sexual diploids of the L. ovalifolium complex, agamospermous tetraploids of the L. binervosum complex and the triploid L. algarvense. In this study, we investigated: (1) cytotype distribution and diversity within and among populations in an overlapping region of diploid and polyploid *Limonium* spp. in south-western Iberia and north-western Morocco; and (2) patterns of geographical parthenogenesis and ecological preferences across a latitudinal gradient on the Atlantic coast. We show here for the first time that L. nydeggeri and L. algarvense are found further south in Morocco than previously reported. Genome size and ploidy estimates showed that the distribution of these species is not random at the overlapping region studied: tetraploid apomicts tend to be found at higher latitudes than the sexual diploids and L. algarvense grows in sympatry at the southern boundaries of the diploids. Natural populations showed a constancy in ploidy in these complexes. However, we report for the first time the occurrence of mixed-ploidy populations of L. ovalifolium s.l., euploid triploids in L. algarvense and aneuploids in the L. binervosum complex. On the Atlantic coasts, L. algarvense followed by L. ovalifolium complexes occur significantly more frequently in thermomediterranean and dry ombrotype habitats than the L. binervosum complex. Significant differences were also observed among taxa in the frequency of occurrences on the most common lithological groups. In conclusion, this work presents the first biogeographical insights for the group based in a coarse-scale analysis of data and it provides evidence of ecological differentiation between the studied Limonium complexes. © 2016 The Linnean Society of London, Botanical Journal of the Linnean Society, 2016

ADDITIONAL KEYWORDS: agamospecies – apomixis – biogeography – chromosome base numbers – cytotypes – ecological characteristics – geographical distribution – habitat – polyploidy.

## INTRODUCTION

Polyploidy, i.e. whole-genome duplications, is an important evolutionary process often associated with changes in the reproductive system of flowering plants (Grant, 1981; Otto, 2007). A major route for

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polyploid emergence in several plant genera is through sexual polyploidization, a process in which polyploids are generated by the formation and fusion of unreduced gametes (Bretagnolle & Thompson, 1995; Ramsey & Schemske, 1998). Hybridization with polyploidy may lead to the establishment of reproductive barriers within a few generations, either via the formation of a diploid  $F_1$  hybrid and subsequent chromosome duplication (allopolyploidy) or via a triploid hybrid bridge, in which a triploid cytotype serves as an intermediate step in the production of a new tetraploid entity (e.g. Ramsey & Schemske, 1998). The shift to apomixis (agamospermy; asexual reproduction via seed formation) is one possible mechanism to avoid hybrid sterility and stabilize these hybrid biotypes (e.g. Asker & Jerling, 1992).

In flowering plants, almost all apomicts are exclusively polyploid (Asker & Jerling, 1992; Carman, 1997), except in a few cases, such as diploid Boechera Á.Löve and D.Löve (Böcher, 1951) and Paspalum L. (Siena et al., 2008). Apomicts are commonly allotetraploid (e.g. Antennaria Gaertn., Bayer, 1997; Potentilla L., Dobeš et al., 2013; Limonium Mill., Palop-Esteban, Segarra-Moragues & González-Candelas, 2007) or autotetraploid (e.g. Paspalum, Hojsgaard et al., 2008; Ranunculus L., Hörandl, 2011), whereas their sexual relatives are usually diploids. Reproductive differentiation and isolation between sexual and apomictic cytotypes is common in natural populations and their distribution is either allopatric (i.e. spatially separated from each other; van Dijk, 2007), or sympatric (i.e. co-occurring in the same population; Elzinga et al., 1987; Talent & Dickinson, 2007; Cosendai & Hörandl, 2010; Dobeš et al., 2013). Additionally, as most apomicts reproduce through facultative apomixis, i.e. the co-existence of apomictic and sexual seed production in the same plant, cytotype diversity can be higher than expected in these populations (e.g. Cosendai & Hörandl, 2010).

Different distributional patterns in related sexual and asexual organisms ('geographical parthenogenesis') have long been reported by several authors (Bierzychudek, 1985; Asker & Jerling, 1992; Kearney, 2005). Apomicts in Taraxacum F.H.Wigg. and Chondrilla L. (van Dijk, 2003), Ranunculus (Hörandl, 2006; Hörandl, Cosendai & Temsch, 2008) and Limonium (Róis et al., 2016) have larger distributions at higher latitudes and elevations than their sexual relatives and tend to colonize previously glaciated areas. Geographical parthenogenesis has been attributed to several factors that may have contributed to replacement of sexuals by apomicts in sympatric areas. In particular, superior colonizing abilities of polyploid apomicts, i.e. the capacity to establish a new population from a single individual or seed (Richards, 2003), have been invoked to explain geographical parthenogenesis (Hörandl et al., 2008). Apomixis also overcomes the need for mating partners after a long-distance dispersal event (Baker, 1967; Hörandl et al., 2008; Pannel et al., 2015). Autonomous apomicts, which do not need pollen at all, could provide an advantage over pseudogamous apomicts that still require fertile pollen for

endosperm formation (Hörandl et al., 2008). For example, in Asteraceae, autonomous apomixis is predominant (Noyes, 2007) and a higher probability of population establishment in remote areas has been correlated with the patterns of geographical parthenogenesis frequently observed in the family (Bierzychudek, 1985; Hörandl et al., 2008). Another theory hypothesizes a connection between geographical parthenogenesis and unidirectional hybridization between apomicts and sexuals which are expected to replace sexual reproduction by apomixis in sympatric areas and is thought to be a causal factor for the wide distribution of apomictic complexes (Mogie, 1992; Mogie, Britton & Stewart-Cox, 2007; Hörandl et al., 2008). Other theories, such as the 'frozen niche-variation' model, suggest that apomictic taxa are adapted to slightly different niche optima and occupy larger and climatically more extreme areas than their closest sexual counterparts (Vrijenhoek & Parker, 2009). An in-depth study of niche differentiation between diploid and tetraploid cytotypes in the alpine Ranunculus kuepferi Greuter & Burdet showed that diploid and tetraploid populations differed in terms of niche optima and breadth, with a shift towards cooler conditions in tetraploids (Kirchheimer et al., 2016). Rather than niche differentiation, it was assumed that a change towards facultative apomixis was decisive for tetraploid establishment, being an effective reproductive strategy to avoid minority cytotype exclusion.

Limonium (sea-lavenders, Plumbaginaceae) is a cosmopolitan species-rich genus of annuals, perennial herbs, shrubs and lianas, often adapted to extreme saline environments (Baker, 1966; Kubitzki, 1993). In this genus, several taxonomically complex groups (herein called 'complexes') have been identified (Erben, 1978, 1993). These are generally characterized by uniparental reproduction (e.g. selffertilization. agamospermy, gynogenesis) and hybridization at some degree among its members, which with polyploidy make it difficult to classify the biodiversity into discrete and unambiguous species (Ennos, French & Hollingsworth, 2005; Cortinhas et al., 2015; Róis et al., 2016). A wide cytological diversity has been reported for this genus, ranging from euploid and aneuploid diploids to octoploids (Erben, 1978, 1993; Arrigoni & Diana, 1993; Georgakopoulou et al., 2006; Castro & Rosselló, 2007; Róis et al., 2012; Cortinhas et al., 2015). In Limo*nium* section *Limonium*, the majority of sexual species are diploid, such as those of L. ovalifolium Kuntze complex (Erben, 1999; Róis et al., 2016), whereas most triploid and tetraploid species, like those in the *Limonium* binervosum (G.E.Sm.) C.E.Salmon complex, are apomicts (Erben, 1978; Ingrouille & Stace, 1985, 1986; Cowan, Ingrouille & Lledó, 1998; Róis *et al.*, 2016). Triploid taxa appear to be highly concentrated in the western Mediterranean region, whereas tetraploid taxa and those with higher ploidy are found along the Atlantic coasts and in the eastern Mediterranean region (Erben, 1978, 1979, 1993; Ingrouille & Stace, 1985, 1986; Brullo & Erben, 1989). A recent study showed that diploid sexual and tetraploid apomictic *Limonium* taxa differed in their latitudinal distributional patterns (Róis *et al.*, 2016). Nonetheless, insights into biogeographical and ecological aspects of sexual and apomictic taxa in *Limonium* in the southern part of the distribution have been largely unexplored.

In this study, we investigated the distributional patterns of diploid sexual species of the L. ovalifolium complex, tetraploid agamospermous species of the L. binervosum complex and triploid L. algarvense Erben. First, we investigated cytotype distribution and diversity within and among populations in south-western Iberia and north-western Morocco, which is known to contain both diploid and polyploid *Limonium* spp. Individuals from natural populations were sampled and DNA ploidy was estimated using flow cytometry; chromosome counts were also made to confirm ploidy estimations. Second, because natural populations showed constancy in ploidy within these complexes, we analysed climatic and geological data available in online databases and the literature for the entire distribution range of the diploid and polyploid *Limonium* complexes studied here. Although a coarse-scale analysis, this constitutes the first biogeographical study for the group and provides the basis for subsequent fine-scale studies in selected populations.

## MATERIAL AND METHODS Study species

The perennial species studied belong to a group of parallel-veined *Limonium* spp., which are distributed in coastal habitats influenced by maritime winds and/or temporary soil submersion by seawater (Róis et al., 2016). Among them are the diploid species of the *L.* ovalifolium complex (2n = 2x = 16 chromosomes), which includes L. ovalifolium Kuntze, L. nydeggeri Erben and L. lanceolatum (Hoffmanns. & Link) Franco (Erben, 1978, 1993, 1999; Róis et al., 2012, 2013), the tetraploid species of the L. binervosum complex (2n = 4x = 32, 35, 36) comprising L. binervosum, L. dodartii Kuntze and L. multiflorum Erben (Erben, 1978, 1993; Ingrouille & Stace, 1986), and *L. algarvense* (2n = 3x = 25) (Erben, 1978, Ingrouille, 1985). Molecular 1993: analyses using methylation-sensitive amplified polymorphism markers and discriminant function analyses of morphometric data using specimens representative of the *L. ovalifolium* and *L. binervosum* complexes resulted in a clear separation within and between groups on the basis of the bract and calyx characteristics (Róis *et al.*, 2013). Moreover, the diploid species in the *L. ovalifolium* complex showed reasonable morphological differentiation and interspecific differences that were more evident when the analysis was focused on the epigenetic variation (Róis *et al.*, 2013).

Populations of the diploid species mostly reproduce by outcrossing, whereas tetraploids reproduce through apomixis (Róis et al., 2016). There is no information on the reproductive system of triploids. Limonium ovalifolium s.s. extends from Atlantic France, through the south-western Iberian Peninsula to Morocco (Erben, 1978, 1993) and L. nydeggeri thrives on the west and south-west coasts of Portugal (Erben, 1999; Espírito-Santo et al., 2012); in contrast, L. lanceolatum is distributed along the northwest to south-west coasts of the Iberian Peninsula and in northern Morocco [Franco, 1984; EUNIS (European Nature Information System), 2014; Fennane, Ibn Tattou & El Oualidi, 2014]. Limonium binervosum s.l. occurs in Britain and Ireland (Curtis & McGough, 1988; Stace, 2010), France (Lahondère & Biorét, 1995, 1996) and the Iberian Peninsula coasts (Pignatti, 1971; Erben, 1978), whereas L. dodartii grows on the French Atlantic coast, in north-western Spain, western Portugal (Erben, 1993; Lahondère & Biorét, 1995, 1996) and in Mediterranean France (Lahondère & Biorét, 1996) and L. multiflorum occurs in western Portugal (Erben, 1978, 1993; Espírito-Santo et al., 2012; Caperta et al., 2014). Finally, the triploid *L. algarvense* is distributed from south-western Portugal to southern Spain (Erben, 1993) and has been reported as having an uncertain presence in Morocco (Fennane et al., 2014).

### PLANT MATERIAL AND FIELD SAMPLING

Thirty-nine populations of the *L. ovalifolium* and *L. binervosum* complexes and *L. algarvense* were sampled in the Iberian Peninsula (Portugal and Spain), Morocco and France. The species were identified using keys from Erben (1993) and herbarium specimens were deposited in the Herbarium Prof. João de Carvalho e Vasconcellos (LISI). Herbarium vouchers representative of all populations and taxa sampled are listed in Appendix 1. All populations were tagged with the Global Positioning System (Appendix 2).

In the field, leaf material from one to 32 individuals was sampled per population (290 individuals in total). Two to three leaves per individual were stored in labelled hermetic plastic bags and maintained at 4 °C until flow cytometric analysis. Additionally, seeds of mother plants of L. algarvense, L. binervosum s.s., L. dodartii and L. lanceolatum were collected from the same natural populations to establish experimental populations under controlled conditions. These seeds were germinated in a growth chamber (Rumed) with a photoperiod of 18 h/6 h of light and dark, respectively, and a temperature of 25 °C until germination (Róis et al., 2012). Seedlings were transferred to jiffy pots and maintained in similar growth conditions. From these individuals, leaves and roots were collected for flow cytometric analyses and chromosome counts, respectively. This procedure enabled us to assign genome sizes unambiguously to each ploidy detected.

### GENOME SIZE AND DNA PLOIDY ESTIMATIONS

Genome size and DNA ploidy were assessed using flow cytometry. Nuclei were isolated following the procedure of Galbraith *et al.* (1983) in which  $0.5 \text{ cm}^2$ of fresh leaf tissue of *Limonium* was chopped with a razor blade, simultaneously with 0.5 cm<sup>2</sup> of fresh leaf tissue of the internal reference standard, in a Petri dish containing 1 mL of WPB buffer (Loureiro et al., 2007). As an internal standard, Pisum sativum 'Ctirad' (2C = 9.09 pg) or Secale cereale 'Dankovské' (2C = 16.19 pg) was used (Doležel *et al.*, 1998). Pisum sativum was used as primary reference standard for most of the samples, with the exception of 18 samples of L. multiflorum and two of L. algarvense, for which S. cereale was used instead. The nuclear suspension was filtered using a 50-µm nylon mesh and 50  $\mu$ g mL<sup>-1</sup> propidium iodide (PI; Fluka) was added to stain the DNA. To avoid staining of double stranded RNA, 50  $\mu$ g mL<sup>-1</sup> of RNAse (Fluka) was also added. After a 5-min incubation period, samples were analysed in a Partec CyFlow Space flow cytometer (532 nm green solid-state laser, operating at 30 mW; Partec). Integral fluorescence and fluorescence height and width emitted from nuclei were collected through a 620-nm band-pass interference filter. After the initial analyses, the amplifier system was set to a constant voltage and gain. Each day, prior to analysis, the instrument stability and linearity were checked either with fluorescent beads or using PI-stained nuclei isolated from P. sativum 'Ctirad'. Results were acquired using Partec FloMax software v2.4d (Partec) in the form of four graphics: histogram of fluorescence pulse integral in linear scale (FL); forward light scatter (FS) vs. side light scatter (SS), both in logarithmic (log) scale; FL vs. time; and FL vs. SS in log scale. For most samples, a polygonal region was defined in the last-named graphic to include only intact nuclei and this region was subsequently used to gate all the other graphics. At least 1300 particles per G<sub>1</sub> peak were analysed per sample (Suda *et al.*, 2007). Genome size estimates were only considered when the CV values of G<sub>1</sub> peaks were < 5%. Samples with higher CV values were discarded and a new sample was prepared. Up to 12 individuals per population were analysed for genome size and additional individuals were analysed using the pooled sample strategy, i.e. three or four individuals were used for ploidy estimation only.

The absolute genome size in mass units (2C in pg; sensu Greilhuber et al., 2005) was obtained using the following equation: Limonium sp. 2C nuclear DNA content (pg) = (Limonium sp.  $G_1$  peak mean/reference standard  $G_1$  peak mean) × genome size of the reference standard. Because genome sizes were obtained for several individuals that were also characterized karyologically (below), DNA ploidy could be inferred for all individuals analysed.

Descriptive statistics were calculated for genome size data (mean, standard deviation of the mean, coefficient of variation and minimum and maximum values). Differences in genome size between ploidies and between species within ploidies were assessed using a one-way ANOVA, followed by a Tukey test for multiple comparisons. For the comparison among ploidies, genome size was  $\log_{10}(x)$ -transformed to achieve normality and homogeneity of variances. A Mann–Whitney U test was used to compare genome size between tetraploids and aneuploids of L. dodar*tii* and between triploids (including individuals with both 24 and 25 chromosomes for which a genome size continuum was observed) and aneuploids of L. algarvense. Statistical analyses were performed using SigmaPlot for Windows v. 12.5 (Systat Software).

#### CHROMOSOME COUNTS

Chromosome counts were made to confirm the ploidy estimated based on genome sizes obtained using flow cytometry. Chromosomes were counted for three plants of each species following the procedure described by Róis et al. (2012). Briefly, root tips were excised and treated with a 2 mm 8-hydroxyquinoline solution for 2 h at 4 °C in the dark and subsequently for 2 h at room temperature to induce c-metaphases. Then, root tips were fixed in a fresh absolute ethanol/glacial acetic acid (3:1) solution overnight and stored in 70% ethanol at -20 °C. Afterwards, root tips were digested in a pectolytic enzyme mixture [2% cellulase (Sigma), 2% cellulase 'Onozuka R-10' (Serva) and 2% pectinase enzyme (Sigma) solution in  $1 \times EB$  (40 mL 0.1 M citric acid-1-hydrate and 60 mL 0.1 M sodium citrate dihydrate; pH 4.8] for 3 h at 37 °C as described by Róis et al. (2012). Chromosome ride (DAPI) (1 mg mL<sup>-1</sup>) in Vectashield (Vector Laboratories). Chromosome preparations were observed using a Zeiss Axioskop 2 fluorescence microscope and photographed with an AxioCam MRc5 digital camera (Zeiss).

### OCCURRENCE DATA COLLECTION AND CHARACTERIZATION

In total, 1776 records were obtained for the L. ovalifolium and L. binervosum complexes from the Global Biodiversity Information Facility (http://www.gbif.org/). These encompassed the entire distribution range of the species from each complex, all over the Atlantic coasts in Europe, namely in Ireland, Britain, France and the Iberian Peninsula, and in Morocco (Baker, 1953; Pignatti, 1971; Erben, 1978, 1993, 1999; Franco, 1984; Ingrouille, 1985; Ingrouille & Stace, 1986; Curtis & McGough, 1988; Lahondère & Biorét, 1995, 1996; Stace, 2010; Espírito-Santo et al., 2012; EUNIS, 2014; Fennane et al., 2014). If no reliable location information could be found for some records, they were excluded from further analysis. Furthermore, erroneous GBIF (Global Biodiversity Information Facility) occurrence points (e.g. unlikely point locations such as those generated by coordinates outside the country border under which the species has been listed; coordinates located inland, and thus in non-coastal areas of the countries; and coordinates falling in the Atlantic Ocean or in the Mediterranean Sea) were removed through visual inspection in the desktop GIS environment. Additional occurrence data based on our own field observations (between 2009 and 2015) across the Iberian Peninsula and Moroccan coasts were added comprising a total of 219 records for both species complexes and L. algarvense. Since occurrence data from the GBIF for some of the species were missing (e.g. L. algarvense, L. lanceolatum, L. nydeggeri), the data analyses were made considering three species groups, i.e. L. ovalifolium complex, L. binervosum complex and L. algarvense. To map the extent of each species distribution, the added occurrence records were georeferenced using Desktop ArcGIS 10.0 (ESRI). After filtering and completion procedures, 618 occurrence records were considered (Supporting Information, Table S1; Fig. 1).

Occurrence data points were then overlapped with bioclimate maps with a grid size of 5000 m (http:// www.globalbioclimatics.org/) to obtain bioclimate (Supporting Information, Table S2) and thermotype and ombrotype (Supporting Information, Table S3) of each location, according to the World Bioclimatic Classification System (Rivas-Martínez, Rivas-Sáenz & Penas, 2011). Additionally, the location points

were characterized by their surface lithology, which was based on information available at the OneGeology Europe Portal with a scale 1:1000 000 (http:// www.onegeology-europe.org/). According to the information gathered on surface lithology, 18 different lithology groups were identified (Supporting Information, Table S4). Each Limonium taxon was characterized by the frequency of each thermotype, ombrotype and lithology group. Differences within and among taxa in the frequencies of climatic and lithological groups were statistically analysed using  $\chi^2$  tests. For the surface lithology, only the four most common lithological groups observed within each Limonium group were considered for statistical purposes. These statistical analyses were carried in R version 3.1.1 (R Development Core Team).

Available information for plant communities of the studied *Limonium* spp. is scarce and incomplete and this therefore led us to compile a summary (Supporting Information, Table S5) mainly discriminated by dominant plant species according to the available literature (Bendaanoun, 1991; Costa *et al.*, 1998, 2012, 2014; Hammada, 2007).

### RESULTS

### CYTOTYPE COMPOSITION

Flow cytometric analyses enabled us to unambiguously assess the ploidy for 403 individuals (290 collected directly in the field and 113 from plantlets germinated from seeds) from 20 populations of the L. ovalifolium complex (L. ovalifolium s.s., L. nydeggeri and L. lanceolatum), ten populations of the L. binervosum complex (L. binervosum s.s., L. dodartii and L. multiflorum) and seven populations of L. algarvense (Table 1; Fig. 2). Generally, the quality of the flow cytometry histograms was good, with the CV values of the  $G_1$  peaks of the sample and standard averaging 4.20 and 2.65%, respectively. Overall, three cytotypes were detected, namely diploids  $(2n = 2x = 3.55 \pm 0.07 \text{ pg/2C}),$ triploids  $(2n = 3x = 5.69 \pm 0.15 \text{ pg/2C})$  and tetraploids (2n = $4x = 7.49 \pm 0.18$  pg/2C) (Fig. 2) and a few aneuploids (resulting from triploid and tetraploid individuals after the loss of some chromosomes). Not surprisingly, significant differences in genome size were observed between ploidies  $(F_{2,198} = 16\ 815.89,$ P < 0.001).

Despite a few exceptions, most species were ploidy homogeneous. The *L. ovalifolium* complex was uniformly diploid (Figs 2A–C, 3A), except in two mixedploidy populations of *L. lanceolatum* detected in the southern Iberian Peninsula (Figs 2D, 3A; Table 1), with individuals being either diploid (Fig. 2C) or triploid (Fig. 2D). Although only five individuals from



**Figure 1.** Distribution of diploid, triploid and tetraploid *Limonium* spp. in the Atlantic and Mediterranean coasts: 618 occurrence records based on data from the GBIF (399 records) and our own field observations (219 records). Diploid species of the *L. ovalifolium* complex include *L. ovalifolium* s.s., *L. nydeggeri* and *L. lanceolatum*; and tetraploid species of the *L. binervosum* complex include *L. binervosum* s.s., *L. dodartii* and *L. multiflorum*.

each of the two populations were analysed, the triploid cytotype was dominant in one of them (Fig. 3A; Table 1).

Analysis of L. algarvense confirmed that it was triploid (Figs 2E, 3B), although possible aneuploids were also detected in the Moroccan populations (Fig. 3B; Table 1), as they had significantly lower genome sizes (Mann–Whitney U = 0.00,  $n_1 = 4$ ,  $n_2 = 63, P < 0.001$ ; ranging from 5.22 to 5.46 pg/2C; Table 1). A high variation in genome size estimates was also observed in this taxon across the sampled area (Supporting Information, Fig. S1), suggesting that its chromosome number is more variable than previously envisaged or that some individuals have suffered DNA loss. The chromosome counts obtained from the Alvor population in Portugal (Fig. 4C, D; Supporting Information, Fig. S1) suggest that individuals with 24 or 25 chromosomes could occur across the sampled area.

Limonium binervosum s.s., L. dodartii and L. multiflorum were all tetraploid (Figs 2F–H, 3C, D), except for a few individuals of L. dodartii from two populations that appear to be composed of an euploid plants with significantly lower genome sizes than the confirmed tetraploid individuals (possible an euploids:  $6.87 \pm 0.10$  pg, 4x:  $7.43 \pm 0.12$  pg; Mann–Whitney U = 0.00,  $n_1 = 3$ ,  $n_2 = 14$ , P < 0.001; Fig. 3C; Table 1).

Differences in genome size between species of each ploidy were also explored, but no significant differences were observed in either the *L. ovalifolium*  complex  $(F_{2,85} = 2.80, P = 0.066)$  or the *L. binervo-sum* group  $(F_{2,41} = 0.35, P = 0.709)$ .

### CHROMOSOME NUMBERS

Our chromosome counts obtained from metaphase spreads of L. lanceolatum revealed, for the first time, 2n = 2x = 16 chromosomes for this species (Fig. 4A). Furthermore, we established differences in ploidy among L. ovalifolium s.s., L. lanceolatum, L. algarvense, L. binervosum s.s. and L. dodartii (Table 1; Fig. 4). For L. lanceolatum and L. ovalifolium s.s. (Fig. 4A, B), only diploid specimens with 2n = 2x = 16chromosomes were found (Table 1). For L. algarvense, most specimens were triploids with 2n = 3x = 25 chromosomes (Fig. 4D), but individuals with 2n = 3x = 24chromosomes (Fig. 4) were also observed. With regard to L. binervosum s.s., all individuals examined had 2n = 4x = 35 chromosomes (Fig. 4E), and L. dodartii individuals had 2n = 4x = 36 chromosomes (Fig. 4F), suggesting chromosome rearrangements.

#### BIOGEOGRAPHY AND ECOLOGY

Our findings showed that the apomictic *L. binervo*sum complex tended to be distributed at higher latitudes  $(37-55^{\circ}N)$  than the sexual *L. ovalifolium* complex  $(34-48^{\circ}N)$ , whereas *L. algarvense* occurs in an area that overlaps these two complexes and the southern boundaries of diploids  $(35-39^{\circ}N)$ . Differences in the frequency of the thermotypes

		Genome size (2C, pg)				DNA ploidy level		Chromosome counts			
Species	Population	Mean	SD	CV	Min.	Max.	Ν	2n	Ν	Observed	Literature
L. ovalifolium											$16^{*,\dagger}$
,	Portugal, Cascais: Cabo Raso	3.53	0.05	1.45	3.44	3.59	5		20		
	Portugal, Sines: Praia da Ilha do	3.52	0.06	1.62	3.44	3.60	5	2x	17		
	Pessegueiro										
	Portugal, Vila do Bispo: Cabo de Sagres	3.57	0.06	1.66	3.47	3.62	4	2x	4		16
	Portugal, Lagos: Luz	3.52	0.04	1.10	3.49	3.57	3	2x	3		16
	Portugal, Portimão: Ferragudo	3.50	0.04	1.26	3.46	3.55	$^{2}$	2x	2	16	
	Morocco, Rabat: Sidi Abed	3.51	0.06	1.71	3.39	3.57	6	2x	16		
L. nydeggeri											15, 16, $17^{*,\ddagger}$
1 00	Portugal, Peniche: Papôa	3.56	0.01	0.19	3.56	3.57	<b>2</b>	2x	2		
	Portugal, Peniche: Baleal	3.57	0.07	2.09	3.49	3.64	<b>2</b>	2x	2		15, 16
	Portugal, Peniche: N. Sr <sup>a</sup> dos Remédios	3.57	0.06	1.80	3.46	3.67	6	2x	6		16
	Portugal, Cascais: Cabo Raso	3.61	0.05	1.33	3.49	3.65	8	2x	8		16
	Portugal, Aljezur: Pontal da Carrapateira	3.54	0.04	1.14	3.47	3.59	6	2x	18		
	Portugal, Vila do Bispo: Cabo São Vicente	3.55	0.10	2.69	3.41	3.74	6	2x	15		
	Morocco, Rabat: Sidi Abed	3.59	0.04	1.06	3.54	3.65	5	2x	25		
L. lanceolatum	Portugal, Setúbal: Pontal dos Musgos	3.47	0.04	1.29	3.39	3.53	10	2x	10		
	Portugal, Odemira: Vila Nova Milfontes	3.57	0.07	1.90	3.45	3.67	10	2x	22	16	
	Portugal, Tavira: sapal do Barril	3.61	0.08	2.10	3.53	3.78	10	2x	24	16	
	Spain, Avamonte: marshes of Isla	3.54	0.03	0.98	3.48	3.57	4	2x	4		
	Cristina	5.38	_	_	_	_	1	3x	1		
	Spain, Cádiz: Toruños	3.53	_	_	_	_	1	2x	1		
		5.61	0.03	0.55	5.56	5.65	4	3x	4		
	Morocco, Assilah: Oued Tahadart	3.58	0.06	1.71	3.47	3.67	10	2x	32		
	Morocco, Larache: Moulay Bousselham	3.58	0.04	1.15	3.53	3.66	5	2x	25		
L. algarvense											$25^{*,\dagger}$
0	Portugal, Aljezur: Praia da Amoreira	5.66	0.09	1.61	5.53	5.79	6	3x	6		
	Portugal, Portimão: Alvor	5.80	0.09	1.56	5.62	5.90	8	3x	8	24, 25	
	Portugal, V. R.St° António: Castro Marim	5.67	0.04	0.75	5.61	5.73	4	3x	4		
	Portugal, V. R.St° António: Ponta da Areia	5.68	0.04	0.62	5.65	5.73	4	3x	4		
	Spain, Huelva: Spit El Rompido	5.86	0.04	0.64	5.82	5.95	8	3x	8		
	Spain, Cádiz: El Tómbolo de Trafalgar	5.89	0.10	1.67	5.73	5.99	6	3x	6		
	Morocco, Larache: Loukkos 1	5.66	0.09	1.61	5.53	5.78	12	3x	12		
		5.41	0.03	0.50	5.39	5.44	<b>2</b>	an.	2		
	Morocco, Larache: Loukkos 2	5.63	0.09	1.64	5.54	5.77	8	3x	14		
		5.46	_	_	_	_	1	an.	1		
	Morocco, Larache: Loukkos 3	5.61	0.06	1.09	5.55	5.72	5	3x	5		
		5.22	_	_	_	_	1	an.	1		
L. binervosum											27, $35^{*,\dagger,\$}$
	France, Brittany: Saint-Benoit-des- Ondes	7.25	0.03	0.45	7.21	7.29	5	4 <i>x</i>	9		
	France, Brittany: Pointe du Grouin	7.26	0.07	1.00	7.17	7.34	5	4x	10		
	Portugal, Aveiro: São Jacinto	7.44	0.12	1.67	7.22	7.64	6	4x	6		
	Portugal, S. Pedro de Moel: Praia Velha	7.50	0.15	1.95	7.27	7.65	7	4x	7	35	

**Table 1.** Genome size estimates, ploidy and chromosome counts obtained in populations of the *Limonium ovalifolium* complex (*L. ovalifolium*, *L. nydeggeri* and *L. lanceolatum*), *L. algarvense* and the *L. binervosum* complex (*L. binervosum*, *L. dodartii* and *L. multiflorum*) sampled in France, Spain, Portugal and Morocco

Table	1.	Continued

		Genome size (2C, pg)				DNA ploidy level		Chromosome counts			
Species	Population	Mean	SD	CV	Min.	Max.	Ν	2n	Ν	Observed	Literature
L. dodartii											35, 36*,†
	Portugal, Sines: Praia da Oliveirinha	7.51	0.03	0.00	7.48	7.54	<b>2</b>	4x	2		
		6.92	0.13	0.02	6.75	7.08	3	an.	3		
	Portugal, Sines: Porto Covo	7.46	0.06	0.84	7.40	7.56	5	4x	5		
	Portugal, Odemira: Cabo Sardão	7.44	0.04	0.54	7.39	7.50	6	4x	6	36	
		6.96	_	_	_	_	1	an.	1		
L. multiflorum											$32, 35, 36^{*,\uparrow,\ddagger}$
	Portugal, Lourinhã: Vale de Frades	7.56	0.04	0.49	7.53	7.61	3	4x	3		
	Portugal, Mafra: Foz do Lizandro	7.52	0.07	0.89	7.42	7.64	6	4x	6		
	Portugal, Cascais: Cabo Raso	7.70	0.17	2.25	7.32	7.87	12	4x	12		

The following data are given for each taxon, population and ploidy: mean, standard deviation of the mean (SD), coefficient of variation (CV, %), and minimum (Min.) and maximum values (Max.) of the holoploid genome size (2C, pg), followed by sample size for genome size estimates (N); DNA ploidy (2n) and respective sample size (N) is also provided. DNA ploidy levels: 2x, diploid; 3x, triploid; 4x, tetraploid. Chromosome counts observed in this study (Observed) and reported in the bibliography are also provided (Literature).

\*Erben (1978).

<sup>†</sup>Erben (1993).

<sup>‡</sup>Róis et al. (2012).

<sup>§</sup>Ingrouille & Stace (1986).



**Figure 2.** Flow cytometric histograms of propidium iodide-stained nuclei of diploid, triploid and tetraploid ploidies of the *Limonium ovalifolium* complex (A–D; *L. ovalifolium* – A, *L. nydeggeri* – B and *L. lanceolatum* – 2x, C; 3x, D), *L. algarvense* (E) and the *L. binervosum* complex (*L. binervosum* – F, *L. dodartii* – G and *L. multiflorum* – H) analysed simultaneously with the internal standard *Pisum sativum*. The genome size (pg/2C) for each species is indicated.

were observed among the three *Limonium* groups (Thermomediterranean:  $\chi^2 = 29.18$ , P < 0.001; Mesomediterranean:  $\chi^2 = 7.40$ , P = 0.025; Mesotemperate:

 $\chi^2 = 123.30, P < 0.001$ ), with the Supratemperate type only being detected in *L. binervosum* (Fig. 5A; Supporting Information, Tables S2, S3). *Limonium* 



**Figure 3.** Geographical locations of diploid and polyploid *Limonium* populations studied in the overlapping region of south-western Iberia and north-western Morocco. A, *Limonium ovalifolium* complex: *L. ovalifolium s.s.*, white squares; *L. nydeggeri*, white triangles; *L. lanceolatum*, white circles. B, *Limonium algarvense*, grey circles. C, *Limonium binervo-sum* complex: *L. binervosum s.s.*, black squares; *L. dodartii*, black triangles; *L. multiflorum*, black circles. Cytotypes are represented by different colours: white, 2x, diploid; grey, 3x, triploid; black, 4x, tetraploid; stripes, an., aneuploid. Mixed-ploidy populations, i.e. populations with more than one ploidy detected, are denoted with a dot within the symbol and the proportion of cytotypes are provided in an inset pie diagram.

algarvense had the highest occurrence in the Thermomediterranean type followed by the *L. ovalifolium* complex, whereas the *L. binervosum* complex had significantly higher occurrences in the Mesotemperate type (P < 0.05). The same patterns were evident when analysing frequencies of thermotypes within each *Limonium* group (*L. algarvense*:  $\chi^2 = 96.15$ , P < 0.001; *L. ovalifolium*:  $\chi^2 = 190.53$ , P < 0.001; *L. binervosum*  $\chi^2 = 402.30$ , P < 0.001).

There were also differences between *Limonium* taxa in the frequency of occurrence records in some ombrotypes, in particular for Dry ( $\chi^2 = 232.25$ , P < 0.001) Subhumid ( $\chi^2 = 6.93$ , P = 0.031) and Humid types ( $\chi^2 = 148.48$ , P < 0.001). No differences were found for the remaining types (Semiarid:  $\chi^2 = 2.53$ , P = 0.111; Hyperhumid:  $\chi^2 = 2.37$ , P = 0.123) (Fig. 5B; Supporting Information, Tables S2, S3). Under the dry ombrotype, *L. algarvense* followed by the *L. ovalifolium* complex had the highest frequencies of records (P < 0.05). The

frequency of records in the semi-humid ombrotype was significantly higher in *L. ovalifolium* and *L. binervosum* complexes than in *L. algarvense* and the frequency of *L. ovalifolium* records in the humid ombrotype (P < 0.05). The same patterns were evident when analysing frequencies of ombrotypes within each *Limonium* group (*L. algarvense*:  $\chi^2 = 107.37$ , P < 0.001; *L. ovalifolium*:  $\chi^2 = 254.60$ , P < 0.001; *L. binervosum*:  $\chi^2 = 443.24$ , P < 0.001).

Significant differences were observed in the frequency of the lithological groups within *Limonium* taxa (*L. algarvense*:  $\chi^2 = 67.72$ , P < 0.001; *L. ovalifolium*:  $\chi^2 = 19.31$ , P < 0.001; *L. binervosum*:  $\chi^2 =$ 9.44, P = 0.024), with *L. algarvense* occurring more frequently on alluvium, *L. ovalifolium* complex on limestone and *L. binervosum* complex on alluvium, limestone and till surfaces in comparison with other classes observed in each group (Fig. 6; Supporting Information, Table S4). Significant differences were

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**Figure 4.** Mitotic metaphase plates of DAPI-stained metaphase spreads from *Limonium* spp. A, *Limonium* lanceolatum (2n = 2x = 16 chromosomes); B, L. ovalifolium (2n = 2x = 16); C, L. algarvense (2n = 3x = 24); D, L. algarvense (2n = 3x = 25); E, L. binervosum (2n = 4x = 35); F, L. dodartii (2n = 4x = 36).



**Figure 5.** Frequency (%) of occurrence records in thermotypes (A) and ombrotypes (B) in *Limonium algarvense* and the *Limonium ovalifolium* and *Limonium binervosum* complexes. Data are based in GBIF occurrence points from GBIF and our own occurrence data. Thermomed., Thermomediterranean; Mesomed., Mesomediterranean; Mesotemp., Mesotemperate; Supratemp., Supratemperate.

also observed in the frequency of the most common surface lithology groups among *Limonium* taxa (alluvium:  $\chi^2 = 69.67$ , P < 0.001; limestone:  $\chi^2 = 20.57$ , P < 0.001; gravel:  $\chi^2 = 25.08$ , P < 0.001; conglomerate:  $\chi^2 = 23.64$ , P < 0.001; mudstone:  $\chi^2 = 19.36$ ,

P < 0.001), with till being observed only in L. binervosum.

Available data in the literature about the floristic composition of each *Limonium* community did not allow a discriminant statistical analysis (Supporting



**Figure 6.** Frequency (%) of occurence records in surface lithology groups in *Limonium algarvense* and the *L. ovali-folium* and *L. binervosum* complexes. Data are based in GBIF occurrence points from GBIF and our own occurrence data.

Information, Table S5). Both *L. ovalifolium* and *L. binervosum* complexes have the same dominant species, i.e. *Frankenia laevis* L. or *Halimione portulacoides* (L.) Aellen (Costa *et al.*, 1998, 2012, 2014). However, in Morocco, other species can be dominant, including *Mesembryanthemum crystallinum* L., which associates with *L. nydeggeri* (our own unpublished observations), or *Sarcocornia perennis* (Mill.) A.J.Scott and *Juncus rigidus* Desf., which associate with *L. lanceolatum* (Bendaanoun, 1991). *Limonium binervosum* and *L. dodartii* communities are generally composed of only a few other species and in the Portuguese coast they are rare in salt marshes and on sea cliffs.

### DISCUSSION

In this study, we report for the first time the presence of *L. nydeggeri* on coastal cliffs and of *L. algarvense* in saltmarshes in Morocco. We provide novel data on the distribution of cytotypes and the diversity of diploid, triploid and tetraploid *Limonium* populations in a contact zone in the southern parts of their distribution range, i.e. south-western Iberia and adjacent north-western Morocco. Furthermore, using occurrence data available from the GBIF and our own data across a latitudinal gradient in the Atlantic coasts, we show ecological differentiation between the *L. ovalifolium* and *L. binervosum* complexes and *L. algarvense*.

## Taxonomy and cytogenetics of the group: what is known so far?

Limonium is remarkably variable as a result of frequent hybridization and polyploidization events and different reproductive strategies (Baker, 1966; Erben, 1978; Costa et al., 1998; Lledó et al., 2005; Róis et al., 2016). In the group of species addressed in this study, recent research has shown that the L. binervosum complex presents fairly close taxonomic relationships among its members (Erben, 1978; Ingrouille, 1985; Ingrouille & Stace, 1985; Róis et al., 2013) and is clearly phenotypically distinct from the species belonging to the L. ovalifolium complex (Róis et al., 2013). These observations are also supported by the published cytogenetic data. The latter complex has been described as being mainly diploid (Erben, 1978, 1999; Róis et al., 2012; results herein), although aneuploid individuals have been detected in natural populations (Róis et al., 2012); the species of the first complex are mostly tetraploid or aneuploid (Erben, 1978; Ingrouille & Stace, 1985; Róis et al., 2012; results herein), although aneuploid triploid individuals (2n = 3x = 27 chromosomes) have also been reported (Ingrouille & Stace, 1986). In contrast, L. algarvense is considered to be a homogeneously triploid taxon (2n = 3x = 25) that hypothetically arose from crosses between L. ovalifolium and L. binervosum (Baker, 1953; Erben, 1978; Ingrouille, 1985).

Overall, we detected cytotype homogeneity in most populations and species, with these being either diploid (L. ovalifolium complex), triploid (L. algarvense) or tetraploid (L. binervosum complex), which is consistent with previous cytogenetic reports noted above (Erben, 1978, 1993; Ingrouille, 1985; Ingrouille & Stace, 1986; Róis *et al.*, 2012). Remarkably, diploid and triploid species are more widespread and occupy the southernmost latitudes compared with the tetraploids, with L. algarvense occupying the southernmost limits of the distribution of the tetraploids and extending beyond the distribution range of the diploids.

Mixed-ploidy populations composed of diploid and triploid individuals were recorded for the first time in L. lanceolatum. Although the detection of triploid individuals is surprising, it is not completely unexpected. In fact, flow cytometric seed screening in L. ovalifolium s.s. and L. nydeggeri populations demonstrated the production of 2C or 3C seeds in variable proportions (Róis et al., 2012). However, none of the karyotyped seedlings, which were able to grow and produce further seeds, was triploid (Róis et al., 2012), suggesting that triploid seeds might have a fitness disadvantage. Indeed, this could be the reason for their apparent absence from the survey of natural populations studied here (which included 68 individuals from nine localities of L. ovalifolium s.s. and L. nydeggeri, and showed that all individuals were diploid). The triploid individuals may have arisen as a result of the fusion of one reduced (x)with one unreduced (2x) L. lanceolatum gamete. It is noted that the related L. ovalifolium s.s. and L. nydeggeri have been reported to produce reduced and unreduced gametes which can form viable pollen grains of various sizes (Róis et al., 2012). Production of unreduced gametes is quite frequent in nature and is one of the main mechanisms of polyploid formation (Bretagnolle & Thompson, 1995; Ramsey & Schemske, 1998; Levin, 2002). Alternatively, triploid individuals could have originated through hybridization between L. lanceolatum with other Limonium spp., although this remains to be tested. Because mixed-ploidy populations were not expected, our sampling within populations was not extensive enough to unravel the proportion of different cytotypes in natural populations; nonetheless, in one of the populations the triploids seemed to be dominant.

A high variation in genome size estimates was observed in *L. algarvense*, suggesting that the occurrence of different chromosome numbers for this species might be more frequent than previously reported and hypothesized. In addition to the generally assumed triploid plants with 2n = 25 chromosomes, we describe here, for the first time, triploid plants with 2n = 24 chromosomes. Future studies should focus on increasing the number of chromosome counts made from different individuals to evaluate how widespread these contrasting chromosome numbers are for this taxon. The few examples of speciation via an euploidy observed so far in plants showed the prevalence of an euploidy in interspecific hybrids and polyploids, which may contribute to the establishment of new karyotypes (De Storme & Mason, 2014).

Several aneuploid individuals have also been reported previously in L. dodartii populations, although none was found in the current work. Chromosome losses or gains have already been detected in other *Limonium* spp. for the same species and within the same population (Dolcher & Pignatti, 1967, 1971; Diana, 1995; Castro & Rosselló, 2007; Róis et al., 2012). This suggests that chromosome rearrangements resulting in dysploidy or non-disjunction during cell division (De Storme & Mason, 2014). Additionally, a connection between aneuploidy and apomixis has been confirmed in polyploid complexes such as Boechera (Kantama et al., 2007), Potentilla (Asker, 1971), Manihot esculenta Crantz (Nassar et al., 2009) and Limonium (D'Amato, 1940, 1949; Róis et al., 2016). Polyploidization might enable novel ecological adaptations and lead to immediate reproductive isolation from the parental plant(s) (Levin, 1975, 2002; Husband, Baldwin & Suda, 2013). In this context, apomixis is considered to be one of the mechanisms by which newly formed cytotypes can persist and coexist with other cytotypes in the same population (van Dijk, 2007; Dobeš et al., 2013). Additionally, a mate-free reproductive strategy such as apomixis might increase the fitness of a newly generated cytotype and enable its fixation and spread beyond the parental populations (Hojsgaard & Hörandl, 2015).

### Geographical parthenogenesis

A recent phylogeographical and reproductive study in the L. ovalifolium and L. binervosum complexes has provided evidence for a pattern of 'geographical parthenogenesis' (Róis et al., 2016). Here, we show that these taxa present cytotype differentiation along the coasts of the Iberian Peninsula and adjacent North Africa, with tetraploid apomicts tending to be distributed at higher latitudes than the diploid sexuals, and with the triploid L. algarvense occupying areas of sympatry at the southern limits of diploids. The L. binervosum complex was shown to be distributed in north, west and south-west Iberia but was never found at the southern limits of diploids and triploids. It is possible that the observed geographical pattern was driven by different ecological preferences, palaeoclimatic patterns of aridity, ocean currents and/or dispersal patterns by biotic and abiotic vectors.

Other biogeographical studies of apomictic taxa and complexes (e.g. Arnica alpina Salisb., Paspalum simplex Morong, Taraxacum officinale F.H.Wigg. aggregate or Ranunculus auricomus L. aggregate) have also shown similar patterns of 'geographical parthenogenesis', with polyploid apomicts having larger distributions at higher elevations/latitudes than their diploid sexual relatives or having colonized areas differentially (Bierzychudek, 1985; Kearney, 2005; Hörandl *et al.*, 2008). For example, Ranunculus kuepferi sexual diploids are confined to the south-western parts of the Alps, whereas tetraploid apomicts dominate in previously glaciated areas and geographically isolated areas (Cosendai & Hörandl, 2010).

### ECOLOGICAL PATTERNS

Our analysis of climatic data across the entire range showed that L. algarvense followed by the L. ovalifolium complex occurred significantly more frequently in the thermomediterranean type and the dry ombrotypes than the L. binervosum complex in the Atlantic coasts, which instead was distributed across a much more variable range of thermotypes and ombrotypes than the first two taxa. Furthermore, significant differences between taxa were observed in the frequency of occurrences on the most common lithological groups, with L. algarvense occurring more frequently on alluvium, the L. ovalifolium complex on limestone and the L. binervosum complex on alluvium, limestone and till sites. Although it is recognized that the analysis was conducted at a coarse scale, these results suggest a wider ecological plasticity and greater use of the habitat spectrum in apomicts than in sexual species. Still, this hypothesis should be tested through more detailed, finer-scaled vegetation and geological analyses, which should also incorporate soil texture, pH, electrical conductivity and other types of data, to clarify more precisely the ecological preferences and plant communities within and between each complex. In Antennaria parlinii Fernald, a dioecious species that exhibits wide variation in sexuality at the population level, habitat differentiation was described for sexual and asexual populations, with the asexuals frequently found in disturbed roadside ditches and fallow fields, in contrast to the sexual populations that preferred wooded habitats (O'Connell & Eckert, 1999). At the microspecies level, differences at the level of the plant community in which species occurred were pronounced in Rubus fruticosus L. (Weber, 1995) and species of Alchemilla L. (Fröhner, 1990). Similarly, habitat differentiation between sexual and apomictic populations was also reported for the alpine species *Ranunculus*  *auricomus*, in which apomicts showed a fairly even partitioning among habitat types in contrast to their sexual relatives (Hörandl & Paun, 2007). A fine-scale study in diploid and tetraploid *R. kuepferi* populations demonstrated different niche optima and breadths between cytotypes, with tetraploids occupying new niche spaces and leaving some of the diploid niche space unoccupied (Kirchheimer *et al.*, 2016).

## CONCLUSIONS

The cytotype distribution and differentiation of Limonium diploids, triploids and tetraploids appeared not to be random along the studied latitudinal gradient. Coarse-scale data analysis of climatic factors and surface lithology data points to a broader ecological range of tetraploids than diploids and triploids. Future studies should be directed towards exploring the role of fine-scale factors, such as local edaphoclimatic conditions and relevés to clarify better habitat preferences of these highly specialized species of saline environments. This work, which offers the first biogeographical approach of this *Limonium* group, supports a geographical parthenogenesis phenomenon and suggests an ecological differentiation between the studied taxa.

## ACKNOWLEDGEMENTS

We wish to thank Ana Paula Paes (ISA/LEAF, Portugal) and Mohamed El Madihi (Rabat University, Morocco) for support with fieldwork, Wanda Viegas (ISA/LEAF) for many insightful discussions, and Hassan Rankou and two anonymous reviewers for helpful comments. The Haut Commissariat aux Eaux et Foréts et a la Lute Contre la Desértification, Morocco, provided a licence for plant collection in Morocco. The Linnean Society of London through the Percy Sladen Memorial Fund provided a grant for fieldwork in the project Taxonomic reassessment of Limonium spp. (Plumbaginaceae) from Morocco Atlantic coast. The Portuguese Research Foundation, Fundação para a Ciência e a Tecnologia (FCT), provided financial support for this work under the Incentivo/AGR/UI0240/2014. FCT project also financed the work of Sílvia Castro (Starting grant IF/01267/2013) and Joana Costa (SFRH/BD/89910/ 2012).

## CONFLICT OF INTEREST

The authors declare that they have no competing interests.

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### SUPPORTING INFORMATION

Additional Supporting Information may be found in the online version of this article:

Figure S1. Genome size variation in natural L. algarvense populations.

**Table S1.** Synthesis of *Limonium* population occurrence records considered in this study. The three distinct species groups (the *Limonium ovalifolium* and *L. binervosum* complexes and *L. algarvense*) are discriminated, as well as the distinct countries studied and the record sources.

Table S2. Synthesis of bioclimates considered in this study (adapted from Rivas-Martínez et al., 2011).

**Table S3.** Synthesis of thermotypes and ombric types considered in this study (adapted from Rivas-Martínez *et al.*, 2011).

**Table S4.** Synthesis of surface lithology groups considered in this study (adapted from http://www.onegeology-europe.org/).

**Table S5.** Synthesis of plant communities with presence of the studied *Limonium* taxa (from the *Limonium* ovalifolium and *L. binervosum* complexes and *L. algarvense*) according to its dominant species (Bendaanoun, 1991; Costa *et al.*, 1998, 2012, 2014; Hammada, 2007).

Species	Voucher	Site location/Country	Collection date	Collectors
L. algarvense	LISI – 876/2012	Aljezur: Praia da Amoreira/Portugal	03/07/2012	A. Caperta, A. P. Paes, S. Murra, S. Róis
	LISI - 264/2010	Portimão: Alvor/Portugal	25/08/2009	D. Espírito-Santo, R. Caraça
	LISI - 678/2014	Vila Real de St° António: Castro Marim/Portugal	15/09/2014	A. Caperta, S. Conceição, A. P. Paes
	LISI - 680/2014	Vila Real de St° António: Ponta da Areia/Portugal	15/09/2014	A. Caperta, S. Conceição, A. P. Paes
	$\mathrm{LISI}-659/2014$	Huelva: Spit El Rompido/Spain	16/09/2014	A. Caperta., S. Conceição, A. Paes, R. Freitas
	LISI – 681/2014	Cádiz: El Tómbolo de Trafalgar/Spain	17/09/2014	A. Caperta., S. Conceição, A. Paes, R. Freitas
	LISI – 721/2015	Larache: Loukkos/Morocco	06/06/2015	Caperta, Rhazi, Paes, El Madihi, Róis
L. binervosum	LISI – 769/2015	Brittany: Pointe du Grouin/France	03/07/2015	P. Arsénio, P. Rodríguez-González
	LISI – 770/2015	Britanny: St. Benoit des Ondes/France	03/07/2015	P. Arsénio, P. Rodríguez-González
	LISI – 1173/2013	Aveiro: São Jacinto/Portugal	26/07/2013	S. Murra, A. Caperta. A. P. Paes, P. Arsénio
	$\mathrm{LISI}-695/2014$	Marinha Grande: São Pedro Moel/Portugal	13/06/2013	A. Caperta. A. Paes. S. Murra
L. dodartii	LISI - 573/2010	Sines: Oliveirinha/Portugal	07/07/2010	A. Caperta, S. Róis, A. P. Paes
	LISI - 574/2010	Sines: Porto Covo/Portugal	07/07/2010	A. Caperta, S. Róis, A. P. Paes
	$\mathrm{LISI}-591/2010$	Odemira: Cabo Sardão/Portugal	23/05/2009	A. Caperta, A. P. Paes

# Appendix 1. List of herbarium specimen vouchers representative of *Limonium* populations sampled for genome size and ploidy studies; sampling location and date are also provided

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Appendix	1.	Continued
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Species	Voucher	Site location/Country	Collection date	Collectors
L. lanceolatum	LISI – 558/2010	Odemira: Vila Nova Milfontes/Portugal	22/05/2009	A. Caperta, A. P. Paes
	LISI – <i>890/2012</i>	Tavira: Sapal do Barril	31/07/2012	A. Caperta, D. Espirito Santo, J. C. Costa
	LISI – 716/2015	Assilah: Oued Tahadart/Morocco	05/06/2015	Caperta, Rhazi, Paes, Madihi, Róis
	LISI – 722/2015	Larache: Moulay Bousselham/Morocco	06/06/2015	Caperta, Rhazi, Paes, El Madihi, Róis
L. multiflorum	LISI – 350/2010	Lourinhã: Vale dos Frades/Portugal	28/06/2009	A. Caperta
	LISI - 1004/2013	Cascais: Cabo Raso/Portugal	12/06/2012	A. Caperta, A. P. Paes
L. nydeggeri	LISI - 891/2012	Peniche: Ilha do Baleal/Portugal	27/07/2012	A. Caperta, A. P. Paes
	LISI – 169/2011	Peniche: N <sup>a</sup> S <sup>a</sup> Remédios/Portugal	13/07/2011	A. Caperta, A. P. Paes, S. Róis, A. Cortinhas
	LISI - 689/2014	Cascais: Cabo Raso/Portugal	12/06/2012	A. Caperta, A. P. Paes
	LISI – 1033/2012	Aljezur: Pontal da Carrapateira/Portugal	04/06/2012	A. Caperta. A. P. Paes. S. Róis. S. Murra
	LISI-254/2010	Vila do Bispo: Cabo S. Vicente/Portugal	15/05/2010	A. Caperta, S. Róis
	LISI - 704/2015	Rabat: Sidi Abed/Morocco	10/06/2015	Caperta, Rhazi, Paes, El Madihi, Róis
L. ovalifolium	LISI – 581/2010	Cascais: Cabo Raso/Portugal	23/06/2010	A. Caperta. S. Róis. A. P. Paes
	LISI – 1009/2013	Sines: Praia da Ilha do Pessegueiro/Portugal	02/07/1982	J. G. Pedro. J. F. Costa
	LISI – 341/2010	Vila do Bispo: Ponta de Sagres/Portugal	23/08/2009	A. Caperta, A. R. Antunes
	LISI - 343/2010	Lagos: Praia da Luz/Portugal	16/10/2009	A. Caperta, A. R. Antunes
	LISI – 536/2010	Ferragudo: Ponta do Altar/Portugal	06/07/2010	A. Caperta, S. Róis, A. P. Paes
	LISI – 702/2015	Rabat: Sidi Abed/Morocco	10/06/2015	Caperta, Rhazi, Paes, Madihi, Róis

Appendix 2. Localities of the studied populations of the *Limonium ovalifolium* complex (*L. ovalifolium* s.s., *L. nydeggeri* and *L. lanceolatum*), *L. algarvense* and of the *L. binervosum* complex (*L. binervosum*, *L. dodartii* and *L. multiflorum*) sampled in France, Spain, Portugal and Morocco

		Geographical coordinates			
Species	Population	Latitude	Longitude		
L. ovalifolium	Portugal, Cascais: Cabo Raso	38°42′6.96″N	9°28′34.44″W		
,	Portugal, Sines: Praia da Ilha do Pessegueiro	37°49′42.71″N	8°47′27.94″W		
	Portugal, Vila do Bispo: Cabo de Sagres	36°59′39.27″N	8°56′55.52″W		
	Portugal, Lagos: Luz	37°5′14.79″N	8°43′44.74″W		
	Portugal, Portimão: Ferragudo	37°6′23.77″N	8°31′10.39″W		
	Morocco, Rabat: Sidi Abed	33°55′02.52″N	6°58′41.44″W		

## Appendix 2. Continued

		Geographical coordinates			
Species	Population	Latitude	Longitude		
L. nydeggeri	Portugal, Peniche: Papôa	39°22′26.87″N	9°22′38.74″W		
1 00	Portugal, Peniche: Baleal	39°22′44.11″N	9°20′27.54″W		
	Portugal, Peniche: N. Sr <sup>a</sup> dos Remédios	39°22′11.66″N	9°23′44.63″W		
	Portugal, Cascais: Cabo Raso	38°42′34.33″N	9°29′9.77″W		
	Portugal, Aljezur: Pontal da Carrapateira	$37^{\circ}11'42.14''N$	8°54′39.97″W		
	Portugal, Vila do Bispo: Cabo São Vicente	$37^{\circ}1'21.40''N$	8°59′47.63″W		
	Morocco, Rabat: Sidi Abed	33°55′02.52″N	6°58′41.44″W		
L. lanceolatum	Portugal, Setúbal: Pontal dos Musgos	38°32′1.53″N	8°46′48.02″W		
	Portugal, Odemira: Vila Nova Milfontes	37°43′39.92″N	8°46′15.35″W		
	Portugal, Tavira: Sapal do Barril	37°5′26.18″N	7°40′21.77″W		
	Spain, Ayamonte: Marshes of Isla Cristina	37°12′5.73″N	7°22′53.73″W		
	Spain, Cádiz: Toruños	36°30′42.74″N	$6^{\circ}14'1.37''W$		
	Morocco, Assilah: Oued Tahadart	35°35′1.80″N	5°59′16.14″W		
	Morocco, Larache: Moulay Bousselham	34°52′24.90″N	6°17′13.62″W		
L. algarvense	Portugal, Aljezur: Praia da Amoreira	37°20′57.51″N	8°50′42.26″W		
-	Portugal, Portimão: Alvor	37°7′36.58″N	8°36′6.91″W		
	Portugal, V. R.St° António: Castro Marim	37°13′10.57″N	7°26′14.75″W		
	Portugal, V. R.St° António: Ponta da Areia	37°10′38.04″N	7°24′34.07″W		
	Spain, Huelva: Split El Rompido	37°11′25.23″N	6°56′35.57″W		
	Spain, Cadiz: Cabo Trafalgar	$36^{\circ}11'0.05''N$	$6^{\circ}2'6.07''W$		
	Morocco, Larache: Loukkos 1	$35^{\circ}11'52.44''N$	$6^{\circ}6'49.56''W$		
	Morocco, Larache: Loukkos 2	35°11′18.12″N	6°7′50.40″W		
	Morocco, Larache: Loukkos 3	35°11′39.54″N	$6^{\circ}7'44.28''W$		
L. binervosum	France, Brittany: Saint-Benoit-des-Ondes	48°37′21.20″N	$1^{\circ}51'13.8''W$		
	France, Brittany: Pointe du Grouin	48°42′34.38″N	$1^{\circ}50'35.0''W$		
	Portugal, Aveiro: São Jacinto	40°40′26.51″N	8°43′17.63″W		
	Portugal, S. Pedro de Moel: Praia Velha	39°46′19.76″N	9°1′36.50″W		
L. dodartii	Portugal, Sines: Praia da Oliveirinha	37°55′25.08″N	$8^{\circ}48'21.53''W$		
	Portugal, Sines: Porto Covo	37°51′15.41″N	8°47′43.64″W		
	Portugal, Odemira: Cabo Sardão	37°35′49.93″N	8°49′3.90″W		
L. multiflorum	Portugal, Lourinhã: Vale de Frades	39°16′35.42″N	9°20′9.02″W		
	Portugal, Mafra: Foz do Lizandro	38°56′29.51″N	9°24′54.84″W		
	Portugal, Cascais: Cabo Raso	38°42′34.33″N	9°29′9.77″W		