



Botanical Journal of the Linnean Society, 2013, 171, 140-153. With 4 figures

Biogeographical history and diversification of *Petunia* and *Calibrachoa* (Solanaceae) in the Neotropical Pampas grassland

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Received 10 February 2012; revised 14 June 2012; accepted for publication 22 June 2012

The Pampas in the southern Neotropics is a vast region with vegetation composed mainly of grasses, and it may be the least-studied ecosystem in southern South America. Contrary to what was thought until recently, this region is heterogeneous and harbours rich biodiversity and many endemic species; however, little is known about the current geographical distribution and evolution of its plants. Here, we present results from phylogeographical studies on two genera typical of open environments (*Petunia* and *Calibrachoa*) that occur in both the Pampas and the high-altitude grasslands of southern Brazil. The rapid radiations of *Petunia* and *Calibrachoa* are examples of how strong selective pressures for different pollinators, coupled with adaptation to edaphic and climatic differences, may drive the diversification of plants in the Pampas. We also discuss factors that could have affected and driven the diversification and speciation of plants in this environment. Further studies, including some focusing on other taxa, are required to characterize the diversification of plant species in this region more accurately. © 2012 The Linnean Society of London, *Botanical Journal of the Linnean Society*, 2013, **171**, 140–153.

ADDITIONAL KEYWORDS: floral diversification – low-altitude grassland – Neotropics – plant speciation – pollinators – South America.

INTRODUCTION

Biodiversity patterns in a biome are the product of a long and complex history of evolutionary trends involving ecological processes and external environmental forces (Rull, 2011). The current patterns of geographical distribution and plant evolution are the result of geological processes and historical events (e.g. Quaternary climatic episodes) that occurred in the species range (Hewitt, 2000; Vargas, 2003; Pielou, 2008).

Several plant phylogeographical studies have examined the biogeographical effects of Quaternary

climatic changes on speciation and intraspecific differentiation. These have been mostly in temperate regions (Aoki et al., 2006; Bauert et al., 2007; Chen et al., 2007, 2008; Fujii, 2007; Li et al., 2008; Ortiz et al., 2008; Latch et al., 2009; Espíndola et al., 2010), and only a limited number of studies have addressed patterns of plant diversification in the Neotropics (Lorenz-Lemke et al., 2005; Speranza et al., 2007; Miller et al., 2008; Antonelli et al., 2009; Collevatti, Rabelo & Vieira, 2009; Palma-Silva et al., 2009; Ramos, De Lemos & Lovato, 2009; Arana et al., 2010; Hoorn et al., 2010; Ireland et al., 2010; Lage-Novaes et al., 2010; Antonelli & Sanmartin, 2011). The biotic consequences of climate change in the Southern Hemisphere have been discussed in relation to the contraction and expansion of the Andean flora (Simpson & Todzia, 1990; Premoli, Kitzberger &

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Veblen, 2000; Pastorino & Gallo, 2002; Pastorino, Gallo & Hattemer, 2004; Muellner et al., 2005; Serrano-Serrano et al., 2010), the formation of the Atlantic forest refugia (Carnaval & Moritz, 2008; Carnaval et al., 2009), the contraction of the Amazonian forest to islands or refugia (Ledru, 1993; Colinvaux et al., 1996; Prance, 1996; Miles, Grainger & Phillips, 2004), the expansion of tropical and subtropical open formations (Behling, 1995, 1997, 2002; Behling & Pillar, 2007; Behling et al., 2007) and the expansion of seasonally dry forests (Pennington, Prado & Pendry, 2000; Pennington et al., 2004); however, there are few studies on the biogeography and diversification patterns of plants in this context in the tropical and temperate plains of southern South American grasslands (Speranza et al., 2007; Solis-Neffa, 2009).

In the highland fields of southern South America (Behling & Pillar, 2007; Lorenz-Lemke *et al.*, 2010), it has been proposed that the main driver of the speciation of grassland species was isolation by distance (allopatric speciation) in the glacial and interglacial periods. When the forest expanded to the south during the warmer periods (interglacials), it isolated grassland populations, thus disrupting gene flow and promoting diversification and speciation. Less is known about the speciation of lowland species in southern South America, specifically in the Pampas. Unfortunately, few studies have examined speciation in the Pampas.

As the vegetation in the Pampas was probably not affected directly by forest expansion and still contains high plant species diversity, the hypothesis to be tested in this study is whether the biodiversity of the Pampas is also a product of allopatry and geographical isolation, or whether other factors, such as different ecological interactions, can explain the diversity observed in lowland grasslands on the southern edge of the Neotropics. To this end, we show results based on molecular data from two genera typical of open environments (*Petunia* Juss. and *Calibrachoa* Cerv., Solanaceae) that occur in both the Pampas and the high-altitude grasslands of southern Brazil.

STUDY AREA

LOCATION

The Pampas are entirely located in the Neotropics, bounded by the Paranense province to the north and the Espinal province to the west and south (Cabrera & Willink, 1980). This region is one of the largest warm grassland areas in the world, covering approximately 500 000 km² between latitudes 29°S and 39°S. It includes the plains of east-central Argentina, the Uruguayan territory and the southern half of Rio Grande do Sul (RS) Brazilian state (Fig. 1; Berretta, 2001; Pallarés, Berretta & Maraschin, 2005). This study is confined to the northern portion, the Uruguayan province (Cabrera & Willink, 1980), which includes the southern half of RS state, the whole of Uruguay and the southern regions of the Santa Fé and Entre Rios provinces in Argentina.

GEOMORPHOLOGY

The Pampas is a geomorphologically complex region. In the west, it is mostly of Quaternary sedimentary origin, whereas, east of the Uruguay River, sediments range from Devonian to Holocene and were deposited in and over the Brazilian Shield. The Uruguayan and Brazilian Pampas have clear geological continuity and rocky outcrops that date from the Proterozoic to the Quaternary eras. These regions contain a great variety of soil types (derived from basalt, granite, sandstone, silt, etc.) and landforms. The western region of Uruguay presents a major geological discontinuity, with changes in soil and physiography (Grela, 2004).

CLIMATE

During the Quaternary, the glacial cycles resulted in cold, dry conditions interrupted by warmer, wet periods. Consequently, there were several pulses of expansion and retraction of grasslands and, concomitantly, advances and retreats of the northern tropical forests. A dry period occurred at the end of the Pleistocene, during the Last Glacial Maximum (LGM) around 18 000 years ago, with a briefer and less severe dry period in the upper Holocene (Iriondo & García, 1993). During the drier periods, pronounced aeolian activity deflated and redeposited large masses of silt and fine sand over most of the lowlands (Iriondo & García, 1993; Panario & Gutierrez, 1999). Since the LGM, the vegetation of the region has oscillated between xerophytic and tropical and subtropical species. Xerophytic species advanced repeatedly to the north-east during the dry and colder periods and retreated to the south-west during the humid and warmer periods (Iriondo & García, 1993; Iriondo, 1999).

At present, the environment of Uruguay is humid. The average temperatures in the warmest and coldest months are approximately 22 °C and 8 °C, respectively. Rainfall exhibits ample seasonal and annual variation and is more abundant in the north (1500 mm on average) than in the south (1000 mm; Berretta, 2001; Roesch *et al.*, 2009).

VEGETATION

In a broad sense, the Pampas is pure grassland, but the region contains several different physiographic

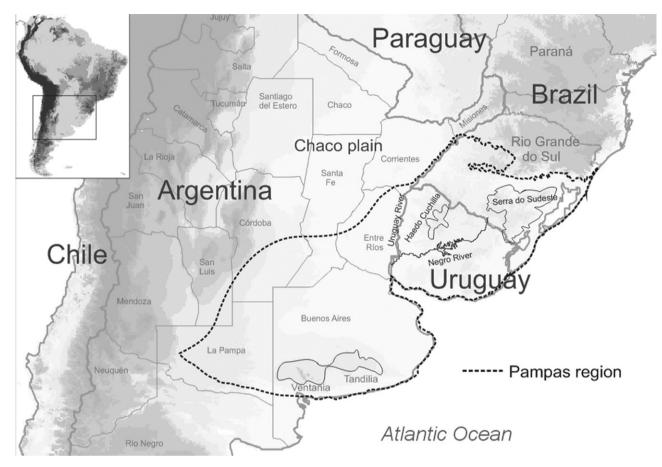


Figure 1. Map representing the extension of the Pampas according to Cabrera & Willink (1980).

formations (Bredenkamp, Spada & Kazmierczak, 2002). In the north, the grassland is invaded by seasonal forests, especially in stream and river valleys. Psammophytic and halophytic steppes occupy the coastal areas, the continental dunes and areas with sandy soils. Shrubby woodlands generally occur on soils with calcareous crusts and consist of shrubland and thornscrub communities, with scarce trees (Quattrocchio et al., 2008). Overbeck et al. (2007) suggested that the southern Brazilian grasslands should be referred to simply as 'campos', which is the most frequently used vernacular term in Uruguay and southern Brazil to refer to natural grazing grasslands covered by grasses, other herbaceous plants, bushes and shrubs, with few trees (Berretta & Nascimento, 1991).

Poaceae are the most species-rich family in the Pampas, with approximately 200 species, including both warm season (C4) and winter (C3) species that are highly characteristic of these grasslands (Berretta, 2001). The most frequently found tribes are Paniceae (including genera with a large number of species, such as *Paspalum L., Panicum L., Axonopus* P.Beauv., *Setaria P.Beauv.* and *Digitaria* Haller), Andropogoneae (with Andropogon L., Bothriochloa Kuntze and Schizachyrium Nees), Eragrosteae (with Eragrostis Wolf and Distichlis Raf.), Chlorideae (with Chloris Sw., Eleusine Gaertn. and Bouteloua Lag.), Poeae (with Bromus L., Poa L., Melica L., Briza L., Lolium L., Dactylis L. and Festuca L.), Stipeae (with Stipa L. and Piptochaetium J.Presl) and Agrostideae (with Calamagrostis Adans and Agrostis L.). Other families that occur in the region include Asteraceae, Fabaceae, Cyperaceae, Apiaceae, Rubiaceae, Plantaginaceae, Oxalidaceae and Solanaceae (Boldrini, 1997; Berretta, 2001).

Anthropogenic land use started approximately 9000 BP (Behling, Pillar & Bauermann, 2005) and, more recently, has had profound impacts on the natural structure of the communities through the introduction of exotic grass species, the conversion of native areas to agricultural lands and grazing areas, and the establishment of *Pinus* L. and *Eucalyptus* L'Her. forests. For example, it is estimated that 50% of the Brazilian Pampas is now composed of vegetation under human management (Roesch *et al.*, 2009). Current human activity in southern Brazil, including extensive areas used as pasture, sometimes makes it difficult to ascertain whether some areas are open because of natural or anthropogenic causes.

PETUNIA AND CALIBRACHOA

Petunia and *Calibrachoa* are included in tribe Petunieae of Solanaceae, members of which are broadly distributed throughout South America. Considering the phylogenetic structure of the genera of Petunieae and their geographical distribution, it has been suggested that *Petunia* and *Calibrachoa* have ancestors of Andean origin (Olmstead *et al.*, 2008).

The species of *Petunia* and *Calibrachoa* were generally considered to be part of the same genus until 1985. Wijsman (1982, 1983) investigated the origin of the garden petunia by crossing different species and, based on these successful crosses, Wijsman & Jong (1985) concluded that the species should be classified into two different groups according to the chromosome numbers and certain morphological characters, such as leaf margins, aestivation, flower symmetry, calyx, anther colour and seed coats, and their classification is supported by cytotaxonomic, reproductive, anatomical and chemical studies (Ellinger *et al.*, 1992; Stehmann *et al.*, 1996; Watanabe *et al.*, 1996a, b; Reis, Sajo & Stehmann, 2002).

The evolutionary history of *Petunia* and *Calibrachoa* has been investigated recently using various molecular tools, and these analyses have revealed short genetic distances among species in each genus, with resultant poorly resolved phylogenetic trees indicating recent diversification (Ando *et al.*, 2005; Kulcheski *et al.*, 2006; Chen *et al.*, 2007). The monophyly of the genera was confirmed, and there is a large genetic distance between the clade *Petunia* plus *Calibrachoa* and other genera in the tribe.

Petunia and Calibrachoa spp. serve as excellent model taxa in which to investigate diversification and speciation in the Pampas. The 14 Petunia spp. are exclusively South American and most are found in southern and south-eastern Brazil (Stehmann et al., 2009) in grasslands, including the Pampas. Known as the garden petunia, this genus has a long history of artificial crosses, including hybrids between P. axillaris (Lam.) Britton, Sterns & Poggenb and P. integrifolia (Hook.) Schinz & Thell, which are disseminated worldwide as ornamental plants (Petunia \times hybrida, Hort. ex Vilm.).

Recent plastid phylogenetic analyses divided *Petunia* into two clades (Kulcheski *et al.*, 2006; Lorenz-Lemke *et al.*, 2010): a highland clade (> 500 m above sea level) with a time to the most recent common ancestor (T_{MRCA}) estimated at ~0.9 Mya (0.6–1.3 Mya) and a lowland clade (species that grow in the Pampas region) with a T_{MRCA} estimated at ~1.1 Mya (0.8– 1.5 Mya). These estimates were obtained from *trnH*- psbA and trnS-trnG plastid spacer sequences and derived by a Bayesian method using a biogeographical calibration point. For both clades, species diversification may have been affected by the climate changes during the glacial and interglacial periods during the Pleistocene (Lorenz-Lemke *et al.*, 2010).

Calibrachoa encompasses 27 species that are found in open areas of southern South America, with an Atlantic subtropical distribution lying between 18°S and 37°S and occurring most densely in the east, along the coast. From the southern boundary, the genus is widely distributed in the Pampas, but is more restricted in the north-west, where rocky and shallow soils support grasslands that extend to the highlands of Santa Catarina and Paraná states. In south-eastern Brazil (the northern limit of the genus distribution in South America), it is represented by a few populations of C. linoides (Sendtn.) Wijsman, a relatively abundant species in the south, and C. elegans (Miers) Stehmann & Semir, an isolated microendemic species found at altitudes > 1000 m in Minas Gerais state. Calibrachoa parviflora (Juss.) D'Arcy is an exception, as it also occurs in North America and Europe. In South America, 15 Calibrachoa spp. inhabit the Paranense biogeographical province and 12 the Pampean and Espinal provinces (according to Cabrera & Willink, 1980).

Most Calibrachoa spp. are self-incompatible and have bee-pollinated flowers (melittophilous). Two exceptions are C. sendtneriana (R.E.Fr.) Stehmann & Semir and C. serrulata (L.B.Sm. & Downs) Stehmann & Semir, which are endemic to the high-altitude grasslands in Santa Catarina and have flowers adapted to hummingbird pollination (Stehmann & Semir, 2001). Calibrachoa parviflora and C. pygmaea (R.E.Fr.) Wijsmann differ substantially from the other species in their reproductive biology and habit. The former is the only self-compatible species in the genus (Tsukamoto et al., 2002), although it has floral traits of the melittophilous syndrome, with extremely small flowers compared with those of other bee-pollinated species. The latter has a unique white hypocrateriform corolla, which is most probably adapted to pollination by hawkmoths. Calibrachoa parviflora and C. pygmaea are also typically herbaceous species with an annual life cycle, whereas other *Calibrachoa* spp. are usually perennial shrubs (Fregonezi et al., 2012).

The melittophilous species have magenta or purple or (infrequently) white or pink corollas. Some taxa have contrasting colours, such as *C. excellens* (R.E.Fr.) Wijsman *atropurpurea* Stehmann & Semir and *C. heterophylla* (Sendtn.) Wijsman, which have a dark purplish ring surrounding the opening of the corolla. In white or pale pink flowers, a striking vein can be observed, as in *C. humilis* (R.E.Fr.) Stehmann & Semir and *C. pubescens* (Spreng.) Stehmann.

MATERIAL AND METHODS

PETUNIA DATA

The sequences of the plastid gene spacers trnH-psbA and trnS-trnG were obtained by Lorenz-Lemke et al. (2006) and Lorenz-Lemke et al. (2010) from lowland species and highland species, respectively. Unpublished sequences from the Petunia integrifolia group (P. integrifolia, P. inflata R.E.Fr., P. interior T.Ando & Hashim. and P. bajeensis T.Ando & Hashim.) of the same genetic markers were included (A.M.C. Ramos-Fregonezi, Universidade Federal do Rio Grande do Sul, Porto Alegre, Brazil, pers. comm.). These sequences were obtained using the same methods as described in the two publications cited above.

The phylogenetic tree for the plastid haplotypes of Petunia spp. was constructed using a Bayesian approach with Beast version 1.6.1 (Drummond & Rambaut, 2007). Two independent runs of 5×10^7 chains were performed, each sampling every 5000 generations. The parameters used were as follows: HKY substitution model with four gamma categories, a Yule tree prior and a relaxed clock model. The software Tracer version 1.5 (available at http:// tree.bio.ed.ac.uk/software/tracer/) was used to check for convergence and adequate effective sample sizes (>200) after the first 10% of generations had been discounted as burn-in. The maximum-clade-credibility trees were estimated using the program TreeAnnotator, which is part of the Beast package. Statistical support for the clades was determined by assessing the Bayesian posterior probability. The haplotype network obtained by Kulcheski et al. (2006) was used to compare those results with this phylogenetic tree and with flower morphology.

CALIBRACHOA DATA

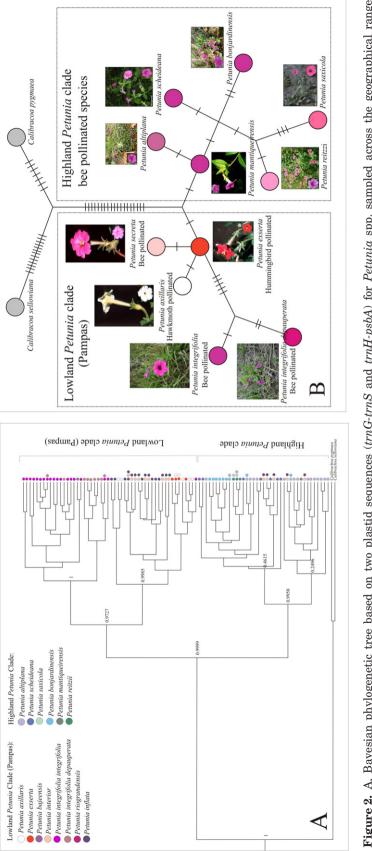
The phylogenetic tree for *Calibrachoa* spp. was based on five noncoding plastid DNA regions: the intergenic spacers *psbB-psbH* and *trnS-trnG* (described by Hamilton, 1999), the intergenic spacer trnL-trnF and the trnL intron (Taberlet et al., 1991) and the intergenic spacer trnH-psbA (Sang, Crawford & Stuessy, 1997). The sequences of the first four regions above were described in a study by Fregonezi et al. (2012), and the trnH-psbA intergenic spacer was added to the dataset to increase the resolution. All samples used were collected in the field, and detailed information about the *trnH-psbA* samples used is given in Supporting Information Appendix S1. Samples of P. axillaris (Lam.) Britton, Sterns & Poggenb. and P. integrifolia were used as the outgroup. The maximum likelihood (ML) phylogenetic analysis was estimated using PAUP* version 4.0b10 (Swofford, 2002) as described in Fregonezi et al. (2012) and in the legend of Figure 3.

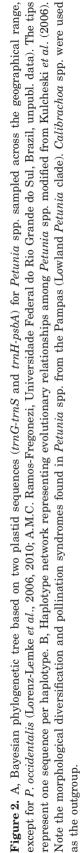
RESULTS AND DISCUSSION

As shown in Figure 2, the lowland (Pampas) Petunia clade comprises species with flowers of different shapes and colours. Petunia axillaris, P. exserta Stehmann and P. secreta Stehmann & Semir have white, red and purple hypocrateriform corollas, respectively, with filaments adnate to the middle of the tube and yellow pollen; P. integrifolia has a purple funnel-form corolla, with filaments adnate to the base of the tube and bluish pollen (Kulcheski et al., 2006; Stehmann et al., 2009). In the Pampas region, these species are often sympatric, with some populations growing together in specific locations (Lorenz-Lemke et al., 2006: personal observations in the field). The plants are self-compatible (except for some lineages of P. ax*illaris*) and the flowers are pollinated by moths, birds and bees, respectively (Ando et al., 1995, 2001; Stehmann et al., 2009; Venail, Dell'Olivo & Kuhlemeier, 2010; Klahre et al., 2011). The highland species have funnel-form or campanulate pink or purple corollas, filaments adnate to the base of the tube and bluish pollen. All highland species have allopatric distributions, are self-incompatible and probably are pollinated by bees (Stehmann et al., 2009; Lorenz-Lemke et al., 2010).

Lorenz-Lemke et al. (2010) suggested that diversification of the highland species occurred as a result of expansion of the ancestral species during glacial periods, followed by fragmentation when forests of Araucaria Juss. spread and surrounded the grassland areas, which subsequently became isolated islands in the areas of highest altitude. This process most probably led to population fragmentation and local differentiation. Which factors might have been important to the speciation of Pampas Petunia (the lowland clade)? Considerable evidence indicates that ecological factors, such as climate and geomorphology, are important to diversification in plant species in the Pampas (see previous studies mentioned above), and may also be important factors in the diversification of lowland Petunia spp. Another question about these species, given the observed variation in reproductive characters (Ando et al., 2005; Lorenz-Lemke et al., 2006), is the role of pollinators; the differences observed among the species could be selected by pollinators and/or by different environmental conditions.

Although the *Petunia* spp. in the Pampas occur in sympatry and may artificially cross-fertilize (Wijsman, 1982; Watanabe *et al.*, 1996a, b; Ando *et al.*, 2001), hybrid forms are rare in nature (Ando *et al.*, 2001; Lorenz-Lemke *et al.*, 2006). Reproductive isolation in the species is thus probably effective. *Petunia hybrida* has been widely used as a model system for molecular genetics, providing a range of genetic and molecular tools (Gerats & Vandenbussche, 2005). These tools





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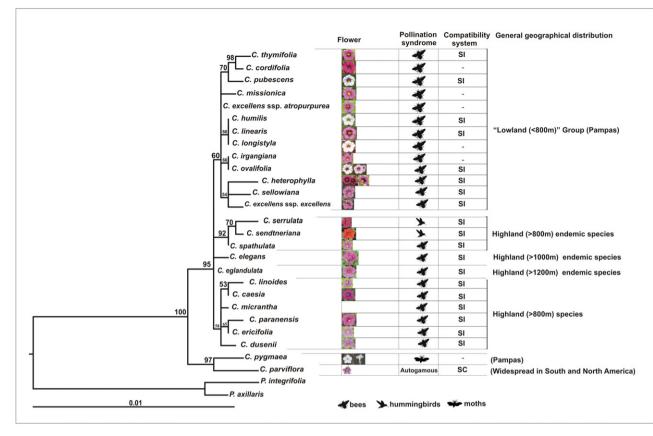


Figure 3. Majority rule consensus tree for the genus *Calibrachoa* of the maximum likelihood analysis from 1000 bootstrap replicates using the Transversional model of nucleotide substitution (TVM) model, based on the combined dataset for the plastid *psbB-psbH*, *trnH-psbA*, *trnL-trnF* and *trnS-trnG* intergenic spacers and the *trnL* intron. The numbers on the branches indicate bootstrap values > 50%. The flowers, floral syndromes and compatibility systems (compiled from Tsukamoto *et al.*, 2002) for each species are indicated, if available.

offer the prospect of an integrated ecological and molecular approach to the analysis of isolation mechanisms in this genus. Previous studies have identified major quantitative trait loci (QTLs) for several traits relevant to pollinator preferences (Stuurman et al., 2004; Galliot et al., 2006; Venail et al., 2010). Dell'Olivo et al. (2011) suggested that the emergence of divergent pollination syndromes alone is unlikely to lead to complete genetic isolation in P. axillaris and P. integrifolia in Uruguay. In addition, gametic isolation is also high in sympatric populations. The floral characters of these species suggest that each conforms to a different pollination syndrome (Gübitz et al., 2009). Petunia *integrifolia* has small, purple, almost scentless flowers that produce hexose-rich nectar (Stuurman et al., 2004; Galliot et al., 2006; Hoballah et al., 2007) and are pollinated by bees (Ando et al., 2001). In contrast, *P. axillaris* flowers are white, have a long corolla tube and produce a strong scent at night; the nectar is sucrose-rich (Stuurman et al., 2004; Oyama-Okubo et al., 2005; Galliot et al., 2006; Hoballah et al., 2007)

and the flowers are pollinated by nocturnal hawkmoths (Ando *et al.*, 2001; Hoballah *et al.*, 2007), although other animals may visit (Ando *et al.*, 2001; our field observations). Although both species occur in sympatry in the wild, natural hybrids have never been found (Ando *et al.*, 2001), suggesting that, in addition to spatial isolation, other strong isolation barriers exist.

Petunia axillaris, P. exserta and P. secreta constitute another example of speciation in the Pampas. They are closely related and share several morphological characteristics, including a long and salverform (hypocrateriform) corolla tube, an erect growth habit and yellow pollen (Fig. 2). The last two species are endemic to the Serra do Sudeste region, RS state, one of the centres of diversity of the genus. Petunia axillaris has a large range throughout the Pampas, and even grows in sympatry with P. exserta in the Serra do Sudeste region (Fig. 1). The major differences among the three species may be associated with different floral syndromes. Petunia exserta has red, nonfragrant flowers with exserted styles and anthers (hummingbird pollination syndrome), whereas P. axillaris has white flowers that are strongly fragrant after dusk (hawkmoth pollination syndrome). Petunia secreta has magenta flowers and melittophily typical of species with very long and narrow tubes; the nectar is virtually inaccessible to bees with short tongues and only pollen is available to visitors. These morphological characteristics suggest that the latter species may have secondarily adjusted melittophily and probably originated from ancestral long-tongued pollinators able to reach the nectar chamber, as in P. axillaris and P. exserta. Observations in a population of P. secreta in RS state showed that this species is visited by bees of the genus Pseudagapostemon (Stehmann, 1999). Petunia axillaris grows in open and sunny habitats, from the Andean slopes in Argentina to Uruguay and southern Brazil, whereas P. exserta can be found only in shady cracks (shelters) in sandstone towers, where it is protected from direct rain and sunlight. Such ecological requirements make P. exserta endemic to specific rock formations in the Serra do Sudeste region of RS state, where it is sympatric with *P. axillaris*, although the latter is found only in open habitats. Lorenz-Lemke et al. (2006) described the occurrence of natural hybrids between P. exserta and P. axillaris; these hybrids are characterized morphologically by pinkish corolla lobes and weakly exserted stamens and styles, and were found only inside shelters, but this phenomenon is not yet well understood.

A. L. A. Segatto (Universidade Federal do Rio Grande do Sul, Porto Alegre, Brazil, unpubl. data) proposed a scenario in which recent and sympatric speciation occurred between *P. axillaris* and *P. ex*serta, suggesting that *P. exserta* may have diverged from a regional population of the widely distributed *P. axillaris* during the LGM and that hybridization has occurred subsequently in some seasons and populations. These species could be an example of sympatric speciation in the Pampas that occurred by adaptive radiation of the floral syndrome and adaptation to different microhabitats.

A similar process may have occurred in the P. axillaris Pampas complex. This species has three recognized subspecies (Ando, 1996): P. axillaris ssp. axillaris (exclusively Pampas), P. axillaris SSD. parodii (Steere) Cabrera (Pampas and Chaco) and P. axillaris ssp. subandina Ando (Pampas and sub-Andean Sierras). These subspecies are geographically delimited on the basis of flower morphology (Ando, 1996; Kokubun et al., 2006). These authors proposed that the Pampas subspecies, P. axillaris axillaris and P. axillaris parodii, are allopatric in Uruguay and that the limit of the distribution of the two subspecies corresponds to the Rio Negro, which could act as a gene flow barrier. In addition, C. Turchetto (Universidade Federal do Rio Grande do Sul, Porto Alegre, Brazil, unpubl. data) identified a major phylogeographical break in the distribution of the subspecies corresponding to climatic discontinuities. Soils could also act as a limiting factor and barrier, as the range of P. axillaris ssp. parodii includes areas on basaltic soil (see description on geology above), suggesting that the environment could affect morphology through selection. Moreover, Venail et al. (2010) studied P. axillaris ssp. axillaris and ssp. parodii, which are similar in all relevant traits for pollinator behaviour, except for floral limb size and tube length. Both of these traits segregate in an interspecific F2 population. Genetically, these traits appear to be complex; three and four medium-effect QTLs were found for limb area and tube length. F2 individuals that differed exclusively in either tube length or limb area were used for pollinator choice experiments. The hawkmoth, Manduca sexta, was more strongly attracted to flowers with larger limb sizes and to flowers with shorter tube lengths. In this sense, the relative importance of pollinator selection could be attributed to the morphology of *P. axillaris*, and our results for the Pampas subspecies of P. axillaris might suggest incipient speciation driven by adaptation to different environmental conditions and pollinatormediated selection (C. Turchetto, unpubl. data).

In summary, studies with *Petunia* spp. are consistent with the hypothesis that historical events, such as climate change in glacial periods, affected the distribution of these species, and that ecological factors, with selective pressure of pollinators on morphological characters, may have been important drivers of speciation.

In relation to *Calibrachoa*, the phylogenetic tree presented in Figure 3 shows a clear division of the genus into two clades: one containing the Pampas species C. parviflora and C. pygmaea and another comprising the remaining species. In the second clade, subclades appear to be associated with the geographical distribution of species. The 'lowland' region refers to the species that are distributed in the Pampas, with specimens collected in the lowlands of RS state, Uruguay and Argentina. In these areas, C. cordifolia Stehmann & L.Aguiar, C. excellens atropurpurea, C. excellens Wijsman excellens, C. heterophylla (Sendtn.) Wijsman, C. humilis, C. irgangiana Stehmann, C. linearis (Hook.) Wijsman, C. longistyla Stehmann & Greppi, C. missionica Stehmann & Semir, C. ovalifolia (Miers) Stehmann & Semir, C. pubescens, C. thymifolia (A.St.-Hil.) Stehmann & Semir, C. parviflora and C. pygmaea can be found.

This phylogenetic tree has short branches and low support values, even using markers with high levels of variation in other studies, including population analysis (Hamilton, Braverman & Soria-Hernanz,

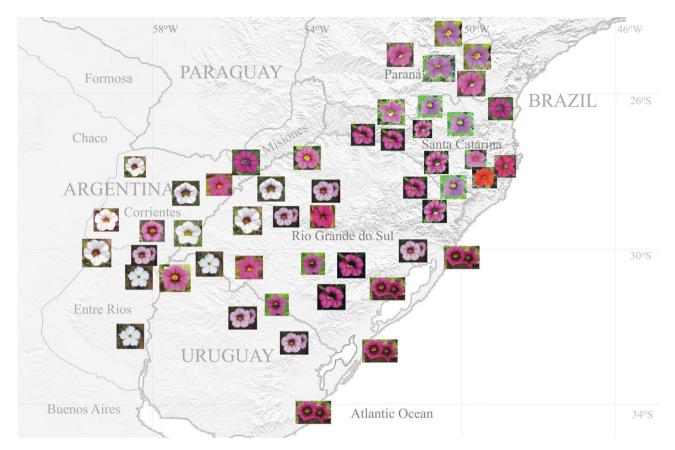


Figure 4. Overall geographical distribution of flower morphological types in *Calibrachoa* spp. The arrangement of figures on the map reflects the general area of occurrence for the representative species of each morphology.

2003; Oh, Kajita & Murata, 2003; Lorenz-Lemke *et al.*, 2006; Bettin *et al.*, 2007). These data suggest rapid morphological divergence in the genus, given the low variation found in the markers used.

The major floral and reproductive divergences probably occurred in the Pampas region. As most species of the genus have melittophilous pollination syndromes, and related genera also have species pollinated by bees, we suggest that the ancestor of the whole group was melittophilous. The Pampas species (also including *C. parviflora* and *C. pygmaea*) have different reproductive strategies, with autogamous species (*C. parviflora*) and self-incompatible species pollinated either by bees or by moths (*C. pygmaea*).

The geographical distribution of flower types in the genus also shows that the highest floral diversity is found in the Pampas (Fig. 4). In the subtropical highlands (the central and northern distribution of the genus), the majority of species have melittophilous syndromes. The exceptions are *C. sendtneriana* and *C. serrulata*, which inhabit the highest mountains in Santa Catarina state and have flowers pollinated by hummingbirds.

Thus, these two main regions (lowland and highland areas of species distribution) appear to have undergone distinct mechanisms of differentiation and speciation in a short period of time. In the highlands, the grasslands were abundant during the colder periods, but were restricted to higher altitudes during the interglacial periods (Behling, 1995, 1997, 2002; Behling & Lichte, 1997). The consecutive cycles of expansion and contraction of grasslands and forests accordingly changed the geographical distribution of *Calibrachoa* populations. The main speciation mechanism therefore appears to be allopatric, without major changes in floral structures.

The Pampas were not strongly affected by the expansion of forest over the grasslands, which have persisted through glacial and interglacial times. The gallery forests found today appeared in the pollen record only approximately 5000 years ago (Behling *et al.*, 2005; Behling & Pillar, 2007), but the number of plant species is as high as that observed in sub-tropical highlands. In this case, the higher speciation rate may be attributed to the type of substrate and to a larger number of pollinators. In the phylogenetic tree (Fig. 3), three groups of closely related species should be highlighted: (1) *C. thymifolia*, *C. cordifolia* and *C. pubescens*, with purple flowers with dark rings, purple and white flowers, respectively; (2)

C. humilis, C. linearis and C. longistyla with white flowers, purple flowers with dark rings and white flowers, respectively; and (3) C. irgangiana and C. ovalifolia, with purple flowers and flowers ranging from pink to white, respectively. Species are found inhabiting mainly sandy soils (e.g. C. heterophylla, C. linearis and C. thymifolia), flooded or wet soils (e.g. C. pygmaea) or latosoils (e.g. C. excellens), with the majority being found on rocky outcrops. Although most melittophilous species have magenta or purple flowers, in the Pampas region, bright flowers (white or pink) and purple flowers with a dark ring are common, suggesting that different pollinators exist in the region. These pollinators may be a mechanism of isolation among groups of sister species. Therefore, in the Pampas region, mechanisms of speciation other than allopatry may have driven the speciation of this group; these could include adaptive evolution to different pollinators. In melittophilous species, the putative pollinators can differ slightly in the way in which

CONCLUSIONS AND FUTURE PROSPECTS

they detect flower colour.

Speciation during grassland formation does not seem to have been affected directly by the expansion and contraction of forests or by ice-sheet advances in relation to Petunia and Calibrachoa lowland clades. In the Pampas region, a climatic gradient and significant soil differences are apparent, and ecological factors were important in the speciation process of these plants. As discussed here, these appear to affect the patterns of diversification and distribution of these species in the Pampas. Studies on other taxa are crucial to the development of a better understanding and, in the future, to the production of a general hypothesis of diversification in this region. Studies of the pollination biology and mating patterns in *Petunia* and *Calibrachoa* spp. and other taxa are essential to the elucidation of the mechanisms of speciation in the Pampas, as pollinators seem to be essential in the evolution of floral traits and in subsequent ecological speciation. In addition to these high-level studies, there is an urgent need for more basic taxonomic studies. Unfortunately, time is running out fast for this area, given the increasing anthropogenic degradation of the biome, with the introduction of exotic plant species as pasture cover and use by the timber industry.

ACKNOWLEDGEMENTS

The authors thank A. P. Lorenz-Lemke, A. M. C. Ramos-Fregonezi, G. Mäder and J. R. Stehmann (Brazil), V. Sollis-Neffa and G. Seijo (Argentina) and P. Speranza (Uruguay) for assistance in field collec-

tion, and two anonymous reviewers and the Associate Editor for criticisms and suggestions that improved the manuscript. This work was supported by the Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq), Coordenação de Aperfeiçoamento de Pessoal de Nível Superior (CAPES) and Fundação de Amparo à Pesquisa do Estado do Rio Grande do Sul (FAPERGS).

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SUPPORTING INFORMATION

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

Appendix S1. Supplementary Material. GenBank accession numbers for *trnH-psbA* sequences used in this study. The information about the collection sites, voucher information and GenBank accession numbers for the other sequences are available in the study of Fregonezi *et al.* (2012).