

The genus *Ruppia* L. (Ruppiaceae) in the Mediterranean region: An overview



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ARTICLE INFO

Article history:

Received 23 December 2013

Received in revised form 17 February 2015

Accepted 19 February 2015

Available online 26 February 2015

Keywords:

Aquatic angiosperms

Ruppia

Shallow waters

Mediterranean region

Overview

ABSTRACT

This paper reviews the current knowledge on the diversity, distribution and ecology of the genus *Ruppia* L. in the Mediterranean region. The genus *Ruppia*, a cosmopolitan aquatic plant complex, is generally restricted to shallow waters such as coastal lagoons and brackish habitats characterized by fine sediments and high salinity fluctuations. In these habitats *Ruppia* meadows play an important structural and functional role. Molecular analyses revealed the presence of 16 haplotypes in the Mediterranean region, one corresponding to *Ruppia maritima* L., and the others to various morphological forms of *Ruppia cirrhosa* (Petagna) Grande, all together referred to as the "*R. cirrhosa* s.l. complex", which also includes *Ruppia drepanensis* Tineo. Populations of hybrid origin between *R. maritima* and *R. cirrhosa* and polyploidy have been detected.

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1. Introduction

The cosmopolitan genus *Ruppia* L. (Ruppiaceae) inhabits a wide variety of shallow systems (coastal lagoons, continental brackish habitats or saltmarsh ponds), where it forms dense and often monospecific meadows that play a key role in the functioning of the ecosystem (Verhoeven, 1979; Calado and Duarte, 2000; Teixeira da Silva and Asmus, 2001; Menéndez, 2002; Malea et al., 2004; Den Hartog and Kuo 2006; Mannino and Sarà, 2006; Pergent et al., 2006; Rodrigues et al., 2009; Obrador and Pretus, 2012). *Ruppia* species are characterized by a simplified morphology that often shows high phenotypic plasticity, leading to taxonomic confusion (Van Vierssen et al., 1981; Hara, 1983; Aedo and Fernández Casado, 1988; Ito et al., 2010). Although the ecology, biomass, productivity and ecophysiology of *Ruppia* species have been widely investigated, the ecotypic and genotypic variation at population level remains only partly understood (Triest and Symoens, 1991; Green and Short, 2003; Den Hartog and Kuo 2006; Triest and Sierens, 2013, 2014).

At present, three species are recognized in the Mediterranean region, *R. maritima* L., *R. cirrhosa* (Petagna) Grande and *R. drepanensis* Tineo (considered by Zhao and Wu, 2008 as synonym of *R.*

cirrhosa), the latter occurring only in the south-western Mediterranean (Comín et al., 1993; Triest and Sierens, 2009a, 2013, 2014; Ito et al., 2013). Recently, hybrid populations between *R. maritima* and *R. cirrhosa* have been detected (Ito et al., 2013; Triest and Sierens, 2013, 2014).

Ruppia taxonomy has been confusing because of the simplified morphology, high phenotypic plasticity, as well as the existence of polyploidy and hybrids that make it difficult to name *Ruppia* species and have a reliable map of their distribution. The present overview, based on studies conducted during the last 40 years, aims at: a) understanding *Ruppia* diversity and mapping its distribution in the Mediterranean basin on the basis of recent molecular studies, b) critically compiling the taxonomic status of Mediterranean *Ruppia*, c) highlighting the main factors affecting *Ruppia* growth, distribution and production, and d) making a comparison with *Ruppia* habitats of other regions in the world.

2. The genus *Ruppia*

2.1. Taxonomy

On the basis of morphological features, the genus *Ruppia* has been traditionally included either within the monogeneric family of Ruppiaceae (Cronquist, 1981; Tomlinson, 1982; Les et al., 1997) or within the family of Potamogetonaceae (Gamerro, 1968; Jacobs

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and Brock, 1982; Dahlgren et al., 1985). Molecular analyses have highlighted a close phylogenetic relationship of Ruppiaceae with seagrass families such as Cymodoceaceae and Posidoniaceae (Les et al., 1997; Zhao and Wu, 2008; Triest and Sierens, 2014).

The high phenotypic plasticity that characterizes this genus has often led to taxonomic confusion (Van Vierssen et al., 1981; Hara, 1983; Aedo and Fernández Casado, 1988; Ito et al., 2010). Disagreement among specialists on the number of *Ruppia* taxa, has led to the proposition of a high number of species, most of which are probably morphotypes of *R. maritima* (Setchell, 1946; Richardson, 1980; Den Hartog, 1981; Kantrud, 1991; Zhao and Wu, 2008), a species with a nearly cosmopolitan distribution (De Oliveira et al., 1983; Orth and Moore, 1988; Malea et al., 2004; Short et al., 2007). Moreover, the occurrence of morphological intermediates with sterile flowers in local populations, suggests that intragenus hybridization may have occurred, making identification even more complicated (Ito et al., 2010).

At present, several species are recognized worldwide: *R. cirrhosa*, *R. maritima*, *R. megacarpa* Mason, *R. tuberosa* Davis & Tomlinson, *R. polycarpa* Mason (Brock, 1982a,b; Zhao and Wu, 2008). *R. maritima* and *R. cirrhosa* are the most widely distributed species, whereas the other three are regarded as endemic to Oceania.

With reference to European habitats, some authors reported only one species, *R. maritima* (Ascherson and Graebner, 1907; Tallon, 1957; Van Ooststroom and Reichgelt, 1964), whereas others recognized two species: *R. cirrhosa* and *R. maritima* (Roze, 1904; Hagström, 1911; Reese, 1962; Den Hartog, 1971; Sfriso, 2010). In the Mediterranean region, three species have been found: *R. maritima*, *R. cirrhosa* and *R. drepanensis*, the latter recognized as an inland ecotype of the SW Mediterranean or a variety, *R. cirrhosa* var. *drepanensis* (Tineo) Symoens (Van Vierssen et al., 1981; Cirujano, 1986; Aedo and Fernández Casado, 1988; Cirujano and García-Murillo, 1990; Triest and Symoens, 1991; Comín et al., 1993; Talavera et al., 1993; Triest and Sierens, 2009a,b, 2010, 2013, 2014). The morpho-anatomy of *Ruppia* species as well as the main morphological characteristics of Mediterranean species are reported in Appendix 1 and Table S1, respectively (see also Figs. 1 and 2). An identification key for the *Ruppia* species, based on morphological characteristics (modified from Zhao and Wu, 2008), is reported in Appendix 2. The key has been modified adding *R. drepanensis* characteristics (Triest and Symoens, 1991; Sfriso, 2008).

2.2. Genetic diversity and chromosome numbers

Only recently, particular attention has been paid to molecular phylogenetics and population genetic diversity (Triest and Symoens, 1991; Triest and Sierens, 2009a,b, 2010, 2013, 2014; Ito et al., 2010, 2013). A first study on isozymes at European and Mediterranean level revealed a cluster of four genetic groups, a low level of polymorphism within *R. maritima* being a separated taxon, and a slight difference between *R. cirrhosa* and *R. drepanensis* (Triest and Symoens, 1991). Isozymes make it difficult to reveal a genetic structure of populations because most populations appear uniform, whereas multiclonal populations are not distinct, probably due to a scarce sampling of enzyme genes (Triest and Symoens, 1991). Similarly, low or absent enzyme polymorphism has been found in seagrasses such as *Zostera* L. (De Heij and Nienhuis, 1992; Williams and Orth, 1998) and *Posidonia* K. Koenig (Capiomont et al., 1996). These aquatic plant groups display rather low allozymic variability, due to the extensive clonal spread (McMillan, 1991; Triest, 1991) and limited hydrophilous pollination (Les, 1988). The lack of genetic variability in genes coding for enzymes, probably lies at the basis of the general difficulties in distinguishing the genetic structure of plant populations from saline habitats such as *Ruppia*, seagrasses and mangrove trees (Triest, 2008). In general,

aquatic plants inhabiting lagoons have very low allozyme diversities (Triest, 1991, 2008). The hypothesis, stated in Triest (2008) (i.e., flowering plants from saline aquatic environments retain lower enzyme polymorphism in their populations, due to constraints on evolutionary transition into the stressful marine environment), should be further tested using DNA polymorphism when coding regions of enzyme genes.

Recently, cpDNA (chloroplast DNA) investigations revealed the presence of 16 haplotypes across the European part of the Mediterranean, one corresponding to *R. maritima* and 15 corresponding to various morphological forms of *R. cirrhosa* all together referred to as the “*R. cirrhosa* s.l. complex”, which includes the *R. drepanensis* (or *R. cirrhosa* var. *drepanensis*) populations (Triest and Sierens, 2010). Those investigations revealed more chloroplast DNA variability at the level of species than would be expected from the traditional view of only two widespread taxa (*R. maritima* and *R. cirrhosa*) and a more restricted taxon, *R. drepanensis* (Triest and Sierens, 2009a,b). *R. maritima* shows a different evolutionary development from other *Ruppia* haplotypes or Mediterranean taxa and might be more rare than previously supposed (Triest and Sierens, 2009a,b). Ito et al. (2010) proposed a world-wide *R. maritima* complex without recognizing the evidence of a distinct *R. maritima* taxon and a polymorphic *R. cirrhosa* complex in Europe (Triest and Sierens, 2009a,b, 2010, 2013, 2014). This hypothesis was updated with phylogenetic analyses of additional Mediterranean samples (Ito et al., 2013) which confirmed the presence of three taxa and hybrids found by Triest and Sierens (2010, 2013). Conclusions about relationships, hybrids and polyploidy are exactly the same for both author groups (Triest and Sierens, 2013, 2014; Ito et al., 2013), but a different name is given to the putative species complex. Triest and Sierens (2009a,b, 2010, 2013, 2014) adhere to the nomenclature of a well-separated diploid *R. maritima* taxon close to an unresolved polymorphic complex of tetraploid *R. cirrhosa* that also includes *R. drepanensis* and traces of hybrid origin. Ito et al. (2013) consider a single world-wide *R. maritima* complex including all phylogenetic clades, even those supported by high bootstrap values. Most of the samples analyzed by Ito et al. (2013) were morphologically identified as *R. maritima*, even outside the clade of the diploid *R. maritima*. It is very unlikely that these world-wide “so-called” *R. maritima* samples (e.g., ‘utahian’, ‘tetraploid alfa’, ‘tetraploid beta’, ‘tetraploid gamma’ and ‘filifolia’ in Fig. 3 of Ito et al., 2013) belong to the same taxon as the type specimen of *R. maritima* from the Baltic sea (‘diploid’ in Fig. 3 of Ito et al., 2013). It may be that the label of *Ruppia* superficially given when a detailed morphological analysis of all plant parts lacks (Den Hartog, personal communication), is a cause of confusion.

The high cpDNA diversity of *Ruppia* in the western Mediterranean basin, representing a center of *Ruppia* diversity, suggests a west–east gradient which can be explained as a historical dissemination of a limited number of haplotypes from the western to the eastern Mediterranean basin (Triest and Sierens, 2010, 2014). CpDNA analyses highlighted the maternal dispersal and spreading through seeds and rhizomes, which for *Ruppia* are more important than dispersal by pollen, whose flow is mainly restricted to a single waterbody as mature pollen floats on the water surface. It has also been proposed that both haplotype recruitment and local dispersal of *Ruppia* propagules by water movement is constrained, or that the genotypes undergo an environmental selection depending on the habitat (Triest and Sierens, 2009a).

The original idea of only two widespread species, *R. cirrhosa* and *R. maritima* (Reese, 1962; Den Hartog, 1971; Dandy, 1980), suggested the presence of a uniform population structure, the separation of which possibly was caused by a long-distance dispersal mediated by birds as generally expected for water plants (Santamaría, 2002). However, cpDNA investigations minimize the effect of strong bird-mediated dispersal (Triest and Sierens,

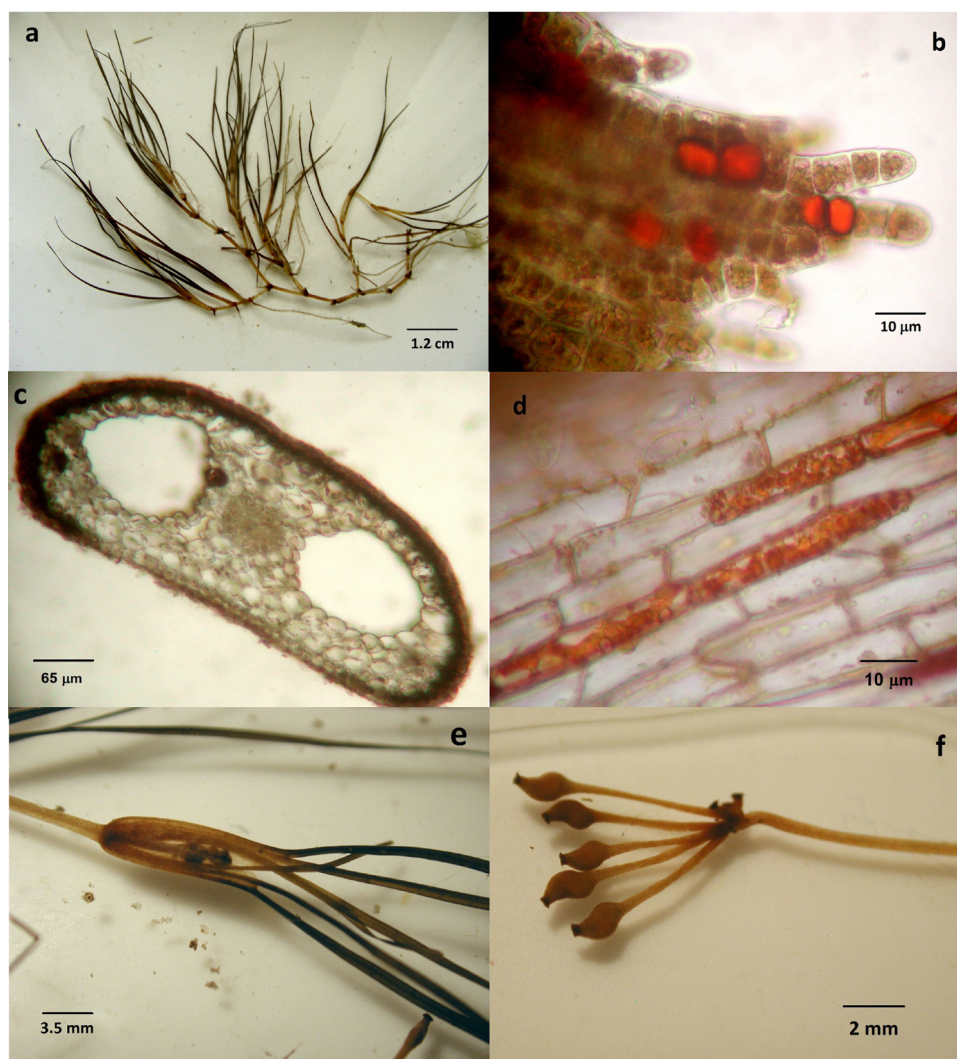


Fig. 1. *Ruppia maritima*: (a) a specimen, (b) a leaf apex, (c) a leaf cross-section, (d) tanniferous cells, (e) a sheathing leaf enclosing the spadix-like inflorescence, (f) a group of achenes.

2009a,b). Moreover, it has been observed that the less fertile haplotypes showed the largest and most continuous distribution (Triest and Sierens, 2013). Therefore, it has been proposed that water movement, or something similar, during and after the last glacial period, determined the distribution of *Ruppia* haplotypes that do not set seeds very frequently in the Mediterranean (e.g., *R. cirrhosa*). In contrast, haplotypes that produce numerous seeds (e.g., *R. drepanensis* and *R. maritima*) were mainly dispersed by bird mediation (Triest and Sierens, 2009b, 2013, 2014).

Differences in chromosome numbers have also been observed. Diploid ($2n=20$) and tetraploid ($2n=40$) races have been reported from several regions of the world, whereas triploids ($2n=30$) and hexaploids ($2n=60$) races have been found only occasionally (Talavera et al., 1993). Previously, the two widespread species, *R. maritima* and *R. cirrhosa*, were considered as diploid and polyploid, respectively (Reese, 1962). However, Van Vierssen et al. (1981) have recognized *R. maritima* as both diploid and tetraploid and *R. cirrhosa* as tetraploid, and Cirujano (1986) considered *R. maritima* as tetraploid and *R. cirrhosa* as diploid. Moreover, Triest and Symoens (1991) recognized diploid and tetraploid entities in both morphological species.

In the Western Mediterranean diploid *R. drepanensis*, tetraploid *R. cirrhosa* and diploid and tetraploid *R. maritima* (Talavera et al.,

1993; Triest and Sierens, 2009a) were recorded. In north-western Spain and in Sicily the diploid *R. maritima* was found (Marchioni Ortu, 1982; Aedo and Fernández Casado, 1988) whereas in Central Spain the tetraploid *R. maritima* and the diploid *R. drepanensis* were recorded (Cirujano, 1986). However, the chromosome counts reported in the literature should be re-evaluated in the context of the nuclear microsatellite and cpDNA haplotypic variation in case of introgressed hybrids (Triest and Sierens, 2014).

The presence of 20 bivalents in *R. maritima* and *R. cirrhosa* at diakinesis would support the hypothesis of an allopolyploid origin for tetraploid races of these species (Talavera et al., 1993).

Moreover, since long arms in large chromosomes do not recombine during meiosis, these regions of the genome might have an adaptive value (Stebbins, 1971; Talavera et al., 1993).

2.3. Distribution

The presence of *R. cirrhosa* has been reported from most countries bordering the Mediterranean Sea where *R. maritima* and *R. drepanensis* are much less common (Table S2).

However, since misidentifications especially of *R. maritima* frequently occur in floristic studies as well as in Genbank (Triest and Sierens, 2014), an accurate identification of *Ruppia* populations is

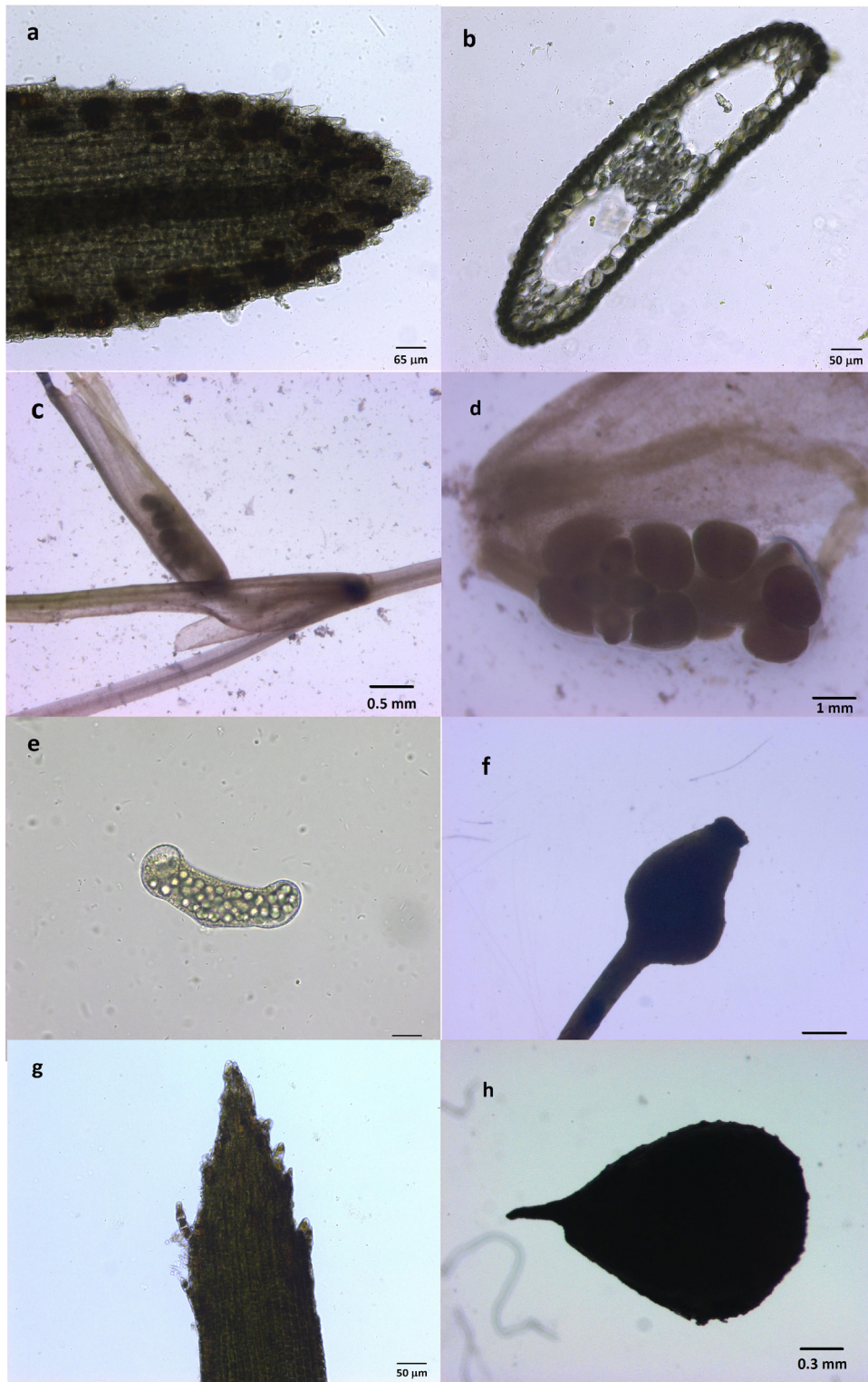


Fig. 2. *Ruppia cirrhosa*: (a) a leaf apex, (b) a leaf cross-section, (c) a sheathing leaf enclosing the spadix-like inflorescence, (d) a flower, (e) a pollen grain, (f) an achen. *Ruppia drepanensis*: (g) a leaf apex, (h) an achen.

needed prior to depict the real distribution of the species. Recently, [Triest and Sierens \(2014\)](#) identified three haplogroup lineages: a *R. cirrhosa* complex (including *R. drepanensis* and ancient hybrid lineages), a *R. maritima* group and some hybrids with most of the diversity occurring in the southern Mediterranean Sea. On the basis of 20 haplotypes, the authors found that the *R. cirrhosa* complex

consists of two common related haplogroups (B and C), a more rare and less related haplogroup A (*R. drepanensis*), an ancient hybrid haplogroup E (consisting of several distant evolutionary lineages) and more recent introgressed hybrids with chloroplast capture. In particular, hybrids between female *R. cirrhosa* and male *R. maritima* or male of *R. cirrhosa* complex at several locations (Southern

Atlantic coast of France, Western Mediterranean coast of Spain and island of Menorca) were observed. A recent northward (Atlantic, North Sea and Baltic coastlines) expansion of *R. cirrhosa* was also observed. The diploid *R. maritima* complex (haplogroup D) was very distinct from the common tetraploid *R. cirrhosa* (Triest and Sierens, 2014).

According to literature, in the extra-Mediterranean areas *R. maritima* seems to be more common than *R. cirrhosa* (Table S3). However, these *R. maritima* populations should be carefully verified. *R. maritima* is the only species recorded in North America (Verhoeven, 1979; Kantrud, 1991; Onuf et al., 2003; Lirman et al., 2008; Cho and Poirrier, 2005; Strazisar et al., 2013), along the South western coasts of Canada (Harrison, 1982) and in India (Rodrigues et al., 2009). In Portugal, in Central and South America and in South Africa, both *R. cirrhosa* and *R. maritima* are present (Póvoa dos Reis, 1973; Verhoeven, 1979; Bortolus et al., 1998; Duarte et al., 2002; Lopez-Calderon et al., 2010; Vromans et al., 2013; Triest and Sierens, 2014). In Australia and New Zealand *R. maritima*, *R. tuberosa*, *R. polycarpa* and *R. megacarpa* are present (Congdon and McComb, 1979; Verhoeven, 1979; Brock, 1982a,b; Jacobs and Brock, 1982; Kim et al., 2013).

Dispersal certainly plays a significant role in determining the distribution of *Ruppia* species. According to Verhoeven (1979), detached vegetative parts of the plants free-float for a long time before landing to the bottom, where they settle rapidly as observed by Orth et al. (1994) for *Zostera marina* L. The same authors also stated that the ripe seeds can be spread by drifting plant parts.

After desiccation, dried plant parts with seeds are dispersed by the wind over considerable distances. Also, several bird species contribute to dispersal. According to Clausen et al. (2002) long-distance dispersal would be rare and the gene flow caused by present day dispersal would play a minor role in comparison to past events. Green et al. (2002) pointed out that, even though only few seeds remain trapped in the intestinal system of birds, they may represent an important dispersal event for clonal plants like *Ruppia*.

2.4. Growth cycle

R. cirrhosa is an annual/perennial species occurring in a wide range of salinity, depth, light and temperature conditions (Verhoeven, 1975, 1979; Cagnoni, 1997; Menéndez, 2002; Menéndez et al., 2002; Agostini et al., 2003; Mannino and Sarà, 2006; Mannino, 2010; Obrador and Pretus 2010; Mannino and Graziano, 2014), in both permanent and temporary lagoons in coastal areas of the NW Mediterranean Sea (Verhoeven, 1979; Gesti, 2000).

After a quiescent period, the length of which is related to temperature, vegetative growth starts with the formation of horizontal rhizomes with monopodial branching. After some time, the horizontal growth changes to vertical, and as soon as flowering begins, branching becomes sympodial (Graves, 1908). Flowering and fructification are more abundant in small water bodies due to the absence of turbulence. *R. cirrhosa* mainly uses “epihydrophilous” (i.e., it takes place at the water surface) cross-pollination, generally less efficient in deep waters (Cook, 1982; Gamarro, 1968; Verhoeven, 1979; Pettit, 1984; Cox, 1988; Cirujano and García-Murillo, 1992; Gesti et al., 2005). Since *R. cirrhosa* brings its flowers up to the water surface, the peduncle length varies according to water depth. In permanent water basins it has a luxuriant vegetative and a moderate generative development. For dispersal, seed transportation by birds and water movement is important. At the end of the vegetative cycle, the shoot tips of *R. cirrhosa* may stop growing vertically and begin to grow horizontally, forming short rhizomes with small roots at each node. These vegetative portions show strong positive geotropism (Verhoeven, 1979; Menéndez et al., 1993).

In temporarily flooded areas, the life cycle must be completed before the basin is dried. Flowering takes place later than in the permanently flooded basins, lasts much shorter, and the production of fruits is higher (Gesti et al., 2005). Also Vromans et al. (2013) observed in warm temperate temporarily estuaries in South Africa the reproductive period of *R. cirrhosa* lasted as long as habitats were permanently flooded and the species only survived the dry phase by seeds.

The growth cycle of *R. maritima* is also significantly affected by environmental conditions. Cho and Poirrier (2005), for example, analyzing some populations from North America observed a positive correlation with high water temperatures. It was also observed that *R. maritima* can grow in unstable intertidal habitats (South America), requiring a higher reproductive resource allocation, and colonize abandoned habitats (i.e., Indian salt pans) (Richardson, 1980; Costa and Seeliger, 1989; Rodrigues et al., 2009).

R. maritima mainly hibernates as seed, but it also displays quiescent vegetative rhizomes in flooded environments. Its generative development is much more luxuriant than that of *R. cirrhosa*, flowering starts earlier and is very abundant. *R. maritima* has a “hypohydrophilous” self-pollination which takes place underwater, but cross-pollination is also possible (Verhoeven, 1979; Richardson, 1983). Under-water (hypohydrophilous) and above-water (epihydrophilous) pollination are both possible in relation to water level changes (Verhoeven, 1979; Lacroix and Kemp, 1997). This pattern fits very well the hypothesis of evolution of underwater pollination from the aerial one (Philbrick, 1988; Philbrick and Les, 1996; Lacroix and Kemp, 1997). This type of pollination is very successful and leads a high seed production, particularly in ephemeral lagoons or estuarine environments (Verhoeven, 1979; Cho and Poirrier, 2005; Kahn and Durako, 2005; Triest and Sierens, 2009a). Desiccation can induce seed dormancy that vice versa can be interrupted by exposure to water (Cho and Sanders, 2009). In temporary habitats, the vegetative parts die and only seed hibernation occurs. Seeds can resist both harsh conditions such as desiccation (Kantrud, 1991; Cho and Sanders, 2009) and even gut-passages of waterfowl and fish, thus allowing a long-distance dispersal (Agami and Waisel, 1988; Figuerola et al., 2002; Charalambidou et al., 2003; Figuerola and Green, 2004). It has been observed that the ingestion of *R. maritima* seeds by birds may positively affect their viability and/or germination rate (Figuerola et al., 2002). However, germination rate increases after ingestion are not necessarily beneficial for the plant, and the final outcome depends on complex interactions with other factors such as herbivore abundance (Figuerola and Green, 2004).

2.5. Ecology

The genus *Ruppia* has a well-defined ecological niche, inhabiting brackish and saline shallow aquatic habitats. Possibly, *Ruppia* is suffering from competition by other water plants outside this brackish range (Verhoeven, 1979; Menéndez and Peñuelas, 1993).

Water level, salinity fluctuations, redox status of surface sediments, wave exposure, self-shading, temperature, desiccation and light availability, with water transparency improving its competitiveness, are generally considered the main factors affecting the distribution and growth of *R. cirrhosa* (Verhoeven, 1979, 1980b; Menéndez and Comín, 1989; Menéndez and Peñuelas, 1993; Adams and Bate, 1994; Santamaría and Hootsmans, 1998; Calado and Duarte, 2000; Azzoni et al., 2001; Menéndez et al., 2002; Gesti et al., 2005; Obrador et al., 2007; Obrador and Pretus, 2008, 2010; Sfriso et al., 2009a; Sfriso, 2010).

In particular, *R. cirrhosa* tolerates a wide range of water temperature (mainly between 5 and 30 °C) and salinity (1.5–60 psu) (Verhoeven, 1979; Menéndez et al., 2002; Malea et al., 2004; Obrador and Pretus, 2010; Mannino and Graziano, 2014). The halo-

tolerance, enhanced by the cytoplasm accumulation of the amino acid proline (Stewart and Lee, 1974; Cavalieri and Huang, 1979; Signorini et al., 2008), makes *R. cirrhosa* more competitive at high salinities (Comín et al., 1991, 1995; Adams and Bate 1994; Murphy et al., 2003; Casagrande and Boudouresque, 2007; Shili et al., 2007; Obrador and Pretus, 2010).

Even though *R. maritima* may survive high salinity fluctuations (Kantrud, 1991), its optimum range is narrower (0.3–15 psu) than that of *R. cirrhosa* (up to 60 psu) (Verhoeven, 1975, 1979). Pulsing salinities may negatively affect *R. maritima* growth rates and reproductive efforts (Verhoeven, 1979; Kantrud, 1991; Bonis et al., 1993; La Peyre and Rowe, 2003). In contrast, populations of *R. maritima* in North America tolerate both low and high salinities quite well (Lirman et al., 2008; Strazisar et al., 2013).

In general, *Ruppia* plants prefer mud and clay sediments, rich in organic material (Giusti and Marsili-Libelli, 2005; Sfriso et al., 2009a; Sfriso, 2010) and have a high resilience (sensu Connell and Sousa, 1983). Eutrophication may negatively affect *Ruppia* survival (Sand-Jensen and Borum, 1991; Sfriso et al., 2007, 2009a,b; Menéndez et al., 1993; Comín et al., 1995; Sfriso and Marcomini, 1996; Viaroli et al., 1996; Azzoni et al., 2001; Teixeira da Silva and Asmus, 2001; Menéndez et al., 2002; Giusti and Marsili-Libelli, 2005; Pergent et al., 2006; Sfriso, 2010), leading to the bed regression or replacement by fast growing thionitrophilous macroalgae, phytoplankton and/or cyanobacteria (Stevenson, 1988; Sand-Jensen and Borum, 1991; Sfriso et al., 2009b).

2.6. Biomass and production

In shallow waters the productivity of macrophytes such as *Ruppia* species plays an important structural and functional role (Mann, 1972; Wetzel, 1975; Sfriso and Facca, 2007; Obrador and Pretus, 2012, 2013) also affecting the carbon cycle (Duarte et al., 2005; Sfriso et al., 2005; Obrador and Pretus, 2012). However, although *R. cirrhosa* productivity is high, its biomass is not significant in the food chain because of the low palatability, due to the presence of several tanniferous cells (Obrador and Pretus, 2012).

R. cirrhosa density (from 4166 to 78,210 shoot m⁻²), biomass (from 12 to 1408 g DW m⁻²) and production (from 36 to 361 g C m⁻² yr⁻¹) have been thoroughly investigated in last decades (Verhoeven, 1980b; Menéndez and Comín, 1989; Menéndez and Peñuelas, 1993; Viaroli et al., 1996; Calado and Duarte, 2000; Duarte et al., 2002; Menéndez et al., 2002; Gesti et al., 2005; Mannino and Sarà, 2006; Obrador et al., 2007; Obrador and Pretus, 2010; Dhib et al., 2013; Vromans et al., 2013; Tables S4 and S5). Most of the *R. cirrhosa* production estimates are based on biomass variations, sometimes including estimates of leaf loss due to mortality during the growing period, as suggested by Menéndez (2002).

Biotic and abiotic external factors, particularly light availability, are important drivers for *R. cirrhosa* production (Calado and Duarte, 2000; Obrador, 2009). Fig. 3 summarizes the relationship between the annual production of *R. cirrhosa* and the depth of the water column for a set of Mediterranean coastal lagoons. A clear negative correlation was observed, suggesting that light is an important limiting factor in determining the annual production and the colonization of shallower bottoms.

The high production of *Ruppia* can lead to carbon limitation (pH > 9) during the summer productivity peaks (Obrador, 2009). The ability to use HCO₃⁻ probably is an important adaptative mechanism for this plant that can also grow in CO₂-depleted waters (Peñuelas and Menéndez, 1990; Hellblom and Axelsson, 2003; Obrador, 2009).

A few data are available on Mediterranean populations of *R. maritima*, while populations from extra-Mediterranean areas have been widely investigated (Gonzalez Gutiérrez, 1977; Howard-Williams and Liptrot, 1980; Verhoeven, 1980b; Flores-Verdugo et al., 1988;

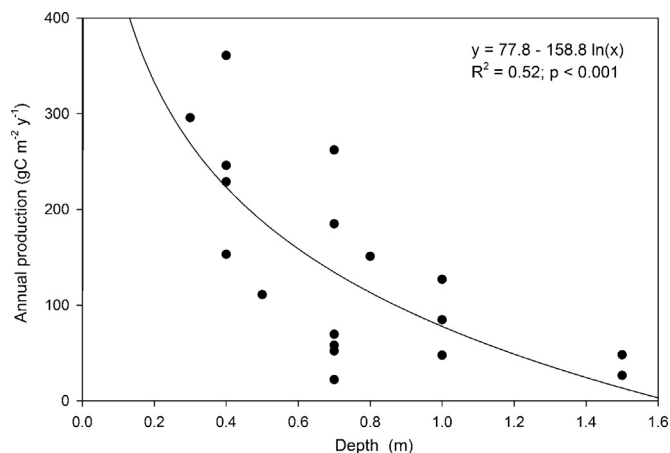


Fig. 3. Annual production of *Ruppia cirrhosa* as a function of water depth from different coastal lagoons. The lagoons corresponding to bibliographic data are Valle Smaracca, Italy (Azzoni et al., 2001; Bartoli et al., 2008); Llacuna de la Tancada, Spain (Menéndez and Comín, 1989; Menéndez, 2002); Albufera des Grau, Spain (Obrador et al., 2007; Obrador and Pretus, 2010); Lake Ichkeul, Tunisia (Casagrande and Boudouresque, 2007); Tvärminne, Finland (Verhoeven, 1980); Camargue, France (Verhoeven, 1980b); Coastal ponds, Netherlands (Verhoeven, 1980b); Certes fishponds, France (Viaroli et al., 1996; Bachelet et al., 2000); Santo André lagoon, Portugal (Calado and Duarte, 2000); Biguglia lagoon, Italy (Pasqualini et al., 2006; Pergent et al., 2006); Sta Giulia lagoon, Italy (Pergent et al., 2006); Fra Ramon, Spain (Gesti et al., 2005).

Cho and Poirrier, 2005; Lopez-Calderon, 2010, Tables S4 and S5). *R. maritima* biomass and productivity are affected by changes of environmental conditions (e.g., temperature, eutrophication, light availability, water movement and basin desiccation), with consequent modifications of local trophic networks, and by grazing (Congdon and McComb, 1979; Verhoeven, 1979, 1980b; Bortolus et al., 1998; Milchakova, 1999; Cho et al., 2009; Lopez-Calderon et al., 2010 and references therein). Moreover, Verhoeven (1980b) observed that the biomass peak of annual populations in temporary brackish habitats (between 25 and 290 g AFDW m⁻²) depends on the time the habitat takes to dry.

Acknowledgements

The authors thank Dr. Orietta Zucchetto for improving the English text and the anonymous reviewers for helpful comments and suggestions. Special thanks go to the Editor for his patience in reviewing the English language and for his critical comments and suggestions that greatly improved the final version of the text.

Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.aquabot.2015.02.005>.

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