REVIEW

Application of biotechnological tools to *Quercus* improvement

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Abstract The genus *Quercus*, which belongs to the family *Fagaceae*, is native to the northern hemisphere and includes deciduous and evergreen species. The trees of the different species are very important from both economic and ecological perspectives. Application of new technological approaches (which span the fields of plant developmental biology, genetic transformation, conservation of elite germplasm and discovery of genes associated with complex multigenic traits) to these long-rotation hardwoods may be of interest for accelerating tree improvement programs. This review provides a summary of the advances made in the application of biotechnological tools to specific oak species. Significant progress has been made in the area of clonal propagation via organogenesis and somatic embryogenesis (SE). Standardized procedures have been developed for micropropagating the most important European (*Q. robur*, *Q. petarea*, *Q. suber*) and American (*Q. alba*, *Q. bicolor*, *Q. rubra*) oaks by axillary shoot growth. Although regenerated plantlets are grown in experimental trials, large scale propagation of oak species has not been carried out.

The induction of SE in oaks from juvenile explants is generally not problematic, although the use of explants other than zygotic embryos is much less efficient. During the last decade, enormous advances have been made in inducing SE from selected adult trees, mainly specimens of pedunculate oak (*Q. robur*) and cork oak (*Q. suber*). Advances in the understanding of the maturation and germination steps are required for better use of embryogenic process in clonal forestry.

Quercus species are late-maturing and late-flowering, exhibit irregular seed set, and produce seeds that are recalcitrant to storage by conventional procedures. Vitrification-based cryopreservation techniques were used successfully in somatic embryos of pedunculate oak and cork oak and an applied genbank of cork oak selected genotypes is now under development. The feasibility of genetic transformation of pedunculate oak and cork oak somatic embryos by means of co-culture techniques with several strains of *Agrobacterium tumefaciens* has also been demonstrated.

To date, most research on the genomics of *Quercus* species has concerned population genetics. Approaches using functional genomics to examine the molecular and cellular mechanisms that control organogenesis and or somatic embryogenesis are still scarce and efforts on the isolation and characterization of genes related to other specific traits should be intensified in the near future, as this would help improve the practical application of clonal forestry in recalcitrant species such as oaks.

Keywords Cryopreservation • genetic transformation • genomics • in vitro tissue culture • micropropagation • oak • somatic embryogenesis • tree improvement

Introduction

Breeding of long-rotation hardwoods remains less well developed than that of short-rotation species (poplars, eucalyptus, willows) and has always lagged behind that of conifers. Investment in projects with an expected lifetime of over 40 years, such as those on long-rotation hardwood species, has never been attractive. *Quercus* species are included in the group of long-rotation hardwoods for which no long-term tree improvement programs have been carried out in either Europe or in North America (Savill and Kanowski 1993; Steiner 1993; Savill et al. 2005).

Conventional tree breeding programs comprises several steps, including: i) provenance trials and or stands of natural origin as the initial phase, ii) seed stand establishment, iii) phenotypic selection of plus trees that may be grafted and planted in clone banks, iv) progeny trials and production of seed orchards for identification of individuals with good genotypes and high combinatory ability, and v) vegetative propagation. Species of the genus *Quercus*, such as *Quercus robur* (pedunculate oak) and *Q. petraea* (sessile oak), usually reach seed-bearing age at between 40 and 100 years old, and the mast seed crops vary according to the individual tree, population region and year (Ducousso and Bordacs 2004), and they are likely to produce good seed crops once every 3-5 years. In addition, vegetative propagation is of extremely limited success because it is very difficult to root cuttings of most tree species (including several *Quercus* species) if they are taken from selected mature trees. Grafting is the method of choice for oak vegetative propagation, but delayed compatibility in *Quercus* has been observed. Expression of graft incompatibility can be delayed up to 7 years and symptoms include significant scion overgrowth of the rootstock, vigorous suckering, and precocious flowering (Coggeshall 1996). According to these limitations in conventional tree breeding approaches, the question remains as to whether biotechnology would be practically useful in the genetic improvement of long-rotation hardwoods and, more specifically, *Quercus* species. Molecular

genetic techniques are undergoing rapid development and may be able to be applied in oak breeding programs, and in physiological and taxonomic studies, as marker traits in selection, identification of individual genotypes, certification of seed orchard products, gene mapping, identification of genes or alleles associated with traits of economic significance (wood properties), etc. (Burley and Kanowski 2005; Grattapaglia et al. 2009). Theoretically, transgenic trees may constitute an important tool for speeding up breeding cycles for traits determined in juvenile woody plants (Flachowski et al. 2009), increasing disease resistance (Fenning 2006), improving wood quality and tree form (Häggman et al. 2006; Grattapaglia et al. 2009), etc. However, there is considerable controversy about the use of genetically modified trees in plantations, so that their use may be restricted in the short- to medium-term future (Bradford et al. 2005; Brunner et al. 2007; Gartland and Oliver 2007; Sederoff 2007; Strauss et al. 2009).

Several recent reviews have described the possibilities of forest biotechnology as an emerging opportunity in relation to tree improvement (Fenning and Gershenzon 2002); some of these reviews cover both coniferous and hardwood species (Nehra et al. 2005) or only hardwoods (Merkle and Nairn 2005; Pijut et al. 2007), but none are devoted specifically to species of the genus *Quercus*. In this review article, we discuss methods that may provide the basis for accelerated improvement of several oak species through biotechnology. Topics addressed in this review include the micropropagation of mature selected material through axillary shoot development, the potential of somatic embryogenesis in light of the most recent results, advances in germplasm conservation through cryopreservation, state-of-the-art in the genetic transformation of the species and the molecular and genomics-related efforts made with the aim of improving these species.

The genus *Quercus*

The genus *Quercus*, which belongs to the family *Fagaceae*, is native to the northern hemisphere and includes deciduous and evergreen species extending from cold latitudes to tropical Asia and the Americas. Its taxonomic classification is very difficult as the genus could be represented by 200 (Neger and Münch 1950), 320 (Krahl-Urban 1959), or 450 species (Krüssmann 1978). The differences in these numbers are partly explained by the definition of interspecific hybrids as separate species and the sub-division of ecological forms into species (Kleinschmit 1993). Frequent hybridization has produced large population of hybrids and high levels of introgression have led to different species in populations sharing up to 50% of their genetic information (Conte et al. 2007; Gömöry and Schmidtová 2007). The genus is subdivided into two subgenera: *Quercus* and *Cyclobalanopsis* and the former is further divided into four sections: *Protobalanus*, *Cerris*, *Quercus* (white oaks) and *Rubrae* (red oaks) (Ducousso and Bordacs 2004).

For the purpose of the present review, we will refer hereafter to the oak species with which most biotechnological approaches have been used.

Twenty-four oak species and hybrids are currently grown in Europe, among which pedunculate oak (*Quercus robur*) and sessile oak (*Q. petraeae*) -which belong to the white oak section- are the most widely distributed and are the most important oak species on the continent, from both economic and ecological perspectives. These oaks, which extend from Ireland to Sicily (Italy) and from the Ural mountains to central Norway (Repo et al. 2008) are large trees that reach 30-40 m in height and live for 800 years or more. The wood from both pedunculate and sessile oaks was used for ship construction and today is employed in furniture making and flooring, timber frame building, veneer production and to make barrels in which wine and spirits are aged. High forest, coppice-with-standards and coppice are the three main silvicultural regimes. Natural regeneration should be a priority but, as it is sometimes difficult, plantations with reproducible material of high genetic value are required (Ducousso and Bordacs 2004).

The European evergreen *Quercus suber* (cork oak) is one of the oak species most characteristic of the Mediterranean ecosystem, and plays important ecological, economic and social roles in parts of southern Portugal and Spain and northern Algeria and Morocco (Knapic et al. 2008) and other countries in the Mediterranean basin. Portugal produces more than half of the cork produced in the world (Catry et al. 2009). This valuable raw material is mainly used to produce wine stoppers (corks).

From the 58 tree-sized native species of oak in the USA, which cover an estimated area of 448,000 km², *Quercus rubra* (red oak) and *Q. alba* (white oak) are the most important and widely distributed (Steiner 1993). Red oak, which is one of the fastest growing of the species, is distributed from northeastern USA reaching west to central Minnesota, eastern Nebraska and Kansas. The geographic range of white oak extends from Maine to Minnesota, south to eastern Texas and northern Florida. The wood of these American oaks is harvested for use in cabinet making, interior finishes, and general construction.

Oak decline syndrome affects oaks worldwide and is of great concern. Oak decline (also known as oak dieback or oak mortality) is an episodic and gradual phenomenon characterized by a general loss of vigor caused by complex interactions between different biotic and abiotic factors and which is not limited to any one species or species group (Wargo et al. 1983; Manion and Lachance 1992; Thomas et al. 2002; Repo et al. 2008). Trees are weakened by environmental stresses such as drought, waterlogging, frost, pests, etc. Oak decline occurs all over Europe and, in its current phase, has been going on since the beginning of the 1980s (Jung et al. 2000). Affected trees display progressive dieback from the tips of the branches, production of chlorotic foliage, and development of epicormic shoots on main branches and stem and

premature autumn leaf colour. Tree mortality is variable and is related to the stand, geographic area, species, etc.

In addition to the episodic phenomenon of oak dieback, there are specific situations where mortality of oaks can only be explained by biotic factors. Fungal pathogens colonizing either the root system (*Armillaria* sp.) or the bole bark (*Biscogniauxia mediterranea*) and bark insects such as *Agrilus* species have been reported as important opportunistic parasites, which are organisms unable to colonize a host unless it has been first weakened as a result of another stress (Marçais and Bréda 2006). The green oak leaf roller (*Tortrix viridiana* L.) is an oligophagus moth that only parasitizes members of the genus *Quercus* and leads to defoliation of oaks in spring (Schroeder and Degen 2008); oak powdery mildew caused by *Erysiphe alphitoides* (formerly *Microsphaera alphitoides*) is the most common disease in the susceptible pedunculate oak (Hajji et al. 2009); the lesions at root or collar level caused by *Phytophthora* sp. are well documented (Robin et al. 1992; Jung et al. 2000; Vettraino et al. 2002). Special mention should be made of sudden oak death, a fungal disease caused by *Phytophthora ramorum*, which has already killed tens of thousands of native coast live oak and tanoak trees in California (Meentemeyer et al. 2008; Brown and Allen-Diaz 2009). This fungus has also been identified in Europe, although the infective capacity of European oaks is, at present, much lower than that of American oaks (Forestry Commission 2009). Nevertheless, expansion of these diseases would increase in the future under predicted conditions of climate change, as expected with *Phytophthora cinnamomi* according to simulation models (Bergot et al. 2004).

In vitro tissue culture

Micropropagation through axillary shoots

Tissue culture techniques were initially applied to different oak species because they provide appropriate tools for rapid production of genotypes to regenerate trees with desired traits and to capture all the genetic superiority without involving any gene segregation. These techniques would, at least theoretically, alleviate the lack of acorn production of selected oak trees on a yearly basis, as well as difficulties in producing offspring with desired traits, difficulties in storing acorns and the low rooting capacity of stem cuttings.

The studies on micropropagation of the most intensively investigated and of the greatest economic importance oak species are summarized in Table 1. The first attempts to micropropagate oak species were carried out, as in many other tree species, with zygotic embryos or seedlings as a source of explants for culture initiation. The disadvantage of using juvenile rather than adult specimens is clear, but was a necessary step to determine the response of oak explants to in vitro conditions. Early studies on the micropropagation of pedunculate and

sessile oaks used embryonic axes as well as shoot apices and nodal segments isolated from 3-6-month-old seedlings (Vieitez et al. 1985; San-José et al. 1990; Chalupa 1993). Although the composition of the mineral media was not decisive in initiating cultures, Woody Plant Medium (WPM) (Lloyd and McCown 1980) and Gresshoff and Doy (1972) medium (GD) were superior to other mineral media for shoot multiplication cultures. In an experiment carried out to identify the most suitable mineral medium in this step in cultures of pedunculate oak, Vieitez et al. (1985) found that, of the eight media tested, the best results were achieved with GD medium. The authors reported that the use of Murashige and Skoog (1962) (MS) medium promoted poorly developed axillary buds, small leaves, no basal callus formation and necrosis that spread from the shoot tips to the whole culture. Cultures grown on MS medium also developed thick, succulent or hyperhydric shoots. The superiority of GD medium was also demonstrated for sessile oak cultures (San-José et al. 1990). The most widely used plant regulator at the multiplication stage is 6-benzylaminopurine (BA) (used at different concentrations), although the effect of thidiazuron in stimulating shoot proliferation was also evaluated (Chalupa 1988). Shoot multiplication in oak cultures does not occur a rosette-like cluster, but by the production of a few, more-or-less elongated shoots. Elongation of shoots may be required prior to rooting. Elongation medium differs from multiplication medium only in that the concentration of BA was either reduced by a factor of ten or was replaced by zeatin (San-José et al. 1988). The addition of zeatin to the multiplication medium improved the proliferation of American oak shoot cultures (Vieitez et al. 1993, 2009).

In the rooting step, isolated shoots (1.5-3 cm) were placed in media containing indole-butyric acid (IBA) or naphthalene-acetic acid (NAA) for a period of 7 or 8 days, with later transfer to an auxin-free medium (Chalupa 1993; Juncker and Favre 1989; Vieitez et al. 1994; Puddephat et al. 1999). Dipping the basal ends of the shoots for 1-2 min in highly concentrated solutions of IBA (1 g/L) and transfer to an auxin-free medium was also carried out (San-José et al. 1988). However, after many experiments, it was found that the best rooting efficiency was achieved by culturing the shoots in media containing 122.5 μ M IBA for 24 h, with subsequent transfer to auxin-free media containing 1% activated charcoal. For all genotypes tested, the charcoal benefited both shoot quality and root system development, and the latter was enhanced by the formation of many lateral roots (Sánchez et al. 1996; Vieitez et al. 2009). In addition, ex vitro rooting experiments were also performed with pedunculate oak microcuttings, which were treated with commercial rooting powder, inserted in peat-perlite substrate and placed under plastic tunnels in a greenhouse (Meier-Dinkel et al. 1993).

While seedling explants are relatively easy to micropropagate, explants from mature trees tend to lose their regeneration potential. When cloning mature trees, it is important to first determine which part of the individual contains the most responsive cells, as some parts of trees

are morphogenetically more competent than others. In hardwood trees and a few gymnosperms, responsive tissues are found at the root-shoot junction, in root or stump sprouts, sphaeroblasts, and epicormic shoots (Bonga et al. 2008, 2010). In vitro plants of pedunculate, sessile, cork and red oaks were regenerated using stump sprouts or epicormic shoots from mature trees, following the above described protocols (Chalupa 1993; Vieitez et al. 1985, 1993, 1994; Favre and Juncker 1987; San-José et al. 1988, 1990; Romano et al., 1992; Evers et al. 1993; Sánchez et al. 1996). These results appear to indicate the good potential for the application of micropropagation techniques in oak forestry. However, several concerns, which may limit application of the technology at a commercial scale, have been identified. The most serious limitations concern: i) the effect of the genotype, ii) the progressive decline of shoot proliferation during successive subcultures and iii) difficulties in propagating mature selected trees lacking stump sprouts or epicormic shoots.

The effect of the genotype on culture initiation and proliferation, which is a common phenomenon in many species, was studied by Meier-Dinkel et al. (1993), who established in vitro cultures from germinated acorns harvested from 11 grafted selected pedunculate oak trees growing in two stands. After 8 months, shoot productivity varied between ten and more than 1000 shoots/genotype; rooting and survival were strongly dependent on the genotype, with values ranging between 10% and 80% for individual genotypes. Furthermore, clonal effects in propagating oak trees were also studied by Juncker and Favre (1989) in pedunculate oak, and data from 16 genotypes derived from seedlings were compared by using principal component and hierarchical cluster analyses. This study confirmed that the between-genotype differences in oak micropropagation are large enough to explain the lack of repeatability in culture establishment, subculture and rooting. No significant between-provenance differences were observed, while the within-provenance differences were high.

The decline in the proliferation capacity of vertically placed oak cultures after several subcultures is a common bottleneck mentioned in most of the publications on the subject. This phenomenon not only affects European oaks but, more specifically, American oaks. All attempts to micropropagate white oak (*Q. alba*) or swamp white oak (*Q. bicolor*) have been unsuccessful. Shoot cultures initiated from terminal and lateral buds of white oak seedlings died through gradual loss of vigor (Schwarz and Schlarbaum 1993). Shoot tip necrosis, dormancy and decline of shoot growth are common problems described as affecting micropropagation of red oak from juvenile seedling material (Vengadesan and Pijut 2007). According to McCown (2000), the decreased capacity for proliferation may be as a result of the episodic growth exhibited by *Quercus*, as the growth is not continuous. The difficulties associated with in vitro culture of oak may be attributed to the inability to achieve uniform and continuous shoot growth. To address the problem, a procedure initially applied to pedunculate oak (San-José et al.

1988; Vieitez et al. 1994) and red oak cultures (Vieitez et al. 1993; Sánchez et al. 1996) was used. Briefly, shoot explants from which the apices (2 mm) have been removed were placed horizontally on the medium and transferred onto fresh medium after 2 weeks, during the 4-week multiplication cycle. After this period, all new shoots longer than 10 mm were harvested and used for multiplication or rooting, and the donor shoots were recultured (transferred to fresh medium) to produce a new crop. The procedure was repeated 3 or 4 times in all (Ballester et al. 2009). By use of this procedure, healthy cultures were produced from pedunculate oak and also from red oak. When the shoots of the American white oaks *Q. alba* and *Q. bicolor* were cultured in the conventional upright position for 4-6 weeks, the strong episodic character of the shoots was shown, with arrested shoot growth generally followed by death of explants. However, culture of the explants in a horizontal position and subsequent reculture of the original explant led, for the first time, to production of vigorous plants of selected American oaks (Vieitez et al. 2009). To explain the beneficial effect of this method in obtaining vigorous shoots, it should be considered that horizontal culture is a form of mechanical stress that may favor synthesis of ethylene in the culture vessel (Anten et al. 2006). It is also possible that horizontal cultures may differ from vertical cultures in both the distribution and transport of endogenous growth regulators and the supply of nutrients.

As mentioned above, oak plantlets have been generated *in vitro* from explants derived from basal shoots or stump sprouts of mature trees, although it was difficult to establish *in vitro* cultures from buds of the current season's growth in the crown (Sánchez 1991). In the absence of mature material with juvenile physiological characteristics, such as stump sprouts or epicormic shoots, rejuvenation or reinvigoration procedures could be applied to mature trees (Ballester et al. 1990). Mature reactive material may be obtained using hedging and stool bed methods, which make it possible to use preformed dormant buds that remain quiescent after early initiation. Outgrowth of dormant buds often leads to the development of physiologically juvenile shoots, not only those located in the lower part of the trunk but also those arising higher up in the tree. Sectioning the trunk of a mature tree to induce epicormic shoots was used as a method of rejuvenation (Evers et al. 1993). However, a variant of this technique (without destroying the tree), consisting of sectioning thick crown branches (3-5 cm) to induce the flushing epicormic shoots, was also used (Vieitez et al. 1994, 2009; Ballester et al. 2009). Briefly, crown branches collected between December and March, cut into 25-30 cm segments, are placed on moist perlite beds in a growth chamber, in order to induce new shoots. The flushed shoots, which develop 10-15 days after severance, are used as source of explants, which develop into shoots exhibiting vigorous growth, long internodes and leaves resembling a more juvenile looking (less lobed) type. After excision of these shoots from the original explants, they may be cultured using the previously described horizontal culture method, and the *in vitro*

multiplication rates recorded were similar to those of cultures of juvenile origin. Following this procedure, genotypes of oak trees older than 300 years have been successfully established in vitro (Vieitez et al. 1994).

After more than 25 years of work on the micropropagation of different oak species through axillary shoot culture, it is possible to conclude the feasibility of the regeneration of whole plants. Interesting technological advances have been carried out that allow oak plants to grow in the field. Plant regeneration by axillary branching is often effective, even with adult material, but it is not sufficiently efficient to be applied on commercial scale. This is in contrast with other genera within the *Fagaceae* family, such as European chestnut, in which large scale micropropagation was reported (Vieitez et al. 2007). Alternative in vitro propagation methods, such as somatic embryogenesis, should be explored.

Somatic embryogenesis

Somatic embryogenesis (SE) is a powerful tool for improvement of forest trees as is considered to be the most appropriate means of in vitro regeneration of woody plants. Large-scale propagation of selected material, genetic transformation, and cryopreservation of elite genotypes are among the most immediate applications of SE. Furthermore, SE may be the only method of regenerating truly juvenile propagules of difficult-to-propagate species. In oaks, induction of the embryogenic process from juvenile explants is generally not problematic, although induction with explants other than mature or immature zygotic embryos is much less efficient. Studies on somatic embryogenesis in different species of the genus *Quercus*, mostly induced from juvenile material, have been summarized in different reviews (Chalupa 1995; Manzanera et al. 1996; Wilhelm 2000), but even since the last date, further progress has been reported in different oak species (Table 2).

Somatic embryo induction According to the definition of recalcitrance in clonal propagation given by Bonga et al. (2010), the most important *Quercus* species belongs to recalcitrant species. The selection of the most reactive explant isolated at the appropriate time applies not only for the propagation through axillary shoot development, but also when the clonal propagation is through SE. As stated in the preceding section, reactive material retaining a certain level of juvenile characters is found in root and stump sprouts, sphaeroblasts, and epicormic shoots of mature oak trees. In the specific case of SE induction, another area of interest involves tissues close to cells that are involved in the sexual process, in which the timing of explant isolation and the application of mild stress are extremely important aspects to be considered.

Expanding leaves excised from epicormic shoots forced to sprout in crown branch segments, according to the methodology previously defined for the micropropagation of mature oak trees via axillary branching (Vieitez et al. 1994), were used as initial explants for the induction of SE in mature oak trees. Following this procedure, the first report on induction of SE in explants of mature *Q. suber* trees was made by Hernández et al. (2001), who initiated cultures from leaves of epicormic shoots, whereas Pinto et al. (2002) also obtained embryogenic calluses in leaves from flushed cuttings of a 60-year-old cork oak tree. Further experiments were carried out for better definition of the induction of SE in cork oak, giving rise to consistent embryogenic systems for the propagation of several hundred-year-old trees (Hernández et al. 2003a). These authors reported that the induction step in leaf explants was performed by successive culture of initial explants in three culture media differing in the concentration of plant growth regulators (PGR): the primary induction medium supplemented with 10 μM BA plus 50 μM NAA, the secondary medium in which the concentration of PGR was reduced to 0.5 μM BA and 0.5 μM NAA, and the expression medium, which is devoid of PGR; the combined application of NAA and BA in primary medium was essential for induction (Hernández et al. 2003a; Toribio et al. 2005). Genotype and time of harvesting, as well as their interaction, significantly influenced the frequency of embryogenic induction, which ranged between 8% and 91% (Hernández et al. 2003b).

Following a similar procedure, SE was also achieved in expanding leaves excised from newly sprouted epicormic shoots forced from branch segments of mature *Q. robur* trees (Toribio et al. 2004). Initiation of somatic embryos was achieved by a multi-stage treatment procedure based on the culture scheme reported for seedling leaf explants of *Q. robur* (Cuenca et al. 1999). The explants were successively cultured in: 1) induction medium (M1) based on MS basal medium and supplemented with 5 mg/l casein hydrolysate, 2.2 μM BA and 21.5 μM NAA, applied in darkness at 25°C for 6-8 weeks; 2) a second medium (M2) of the same composition as the induction medium except that BA and NAA were reduced to respectively 0.44 and 0.54 μM , with culture continued for 4 weeks under a 16 h photoperiod, and 3) expression medium (M3) with the same components but without PGR. The explants were maintained on this medium for 8-10 weeks, for a total 20 weeks after the start of culture. Induction of somatic embryos was correlated with the asymmetric distribution of auxin in the tissues, which appears to be necessary for de novo initiation of organ primordia under culture conditions (Zhao et al. 2008). Accordingly, the next steps in the process were the transfer of oak explants to media with decreasing amounts of PGR, or without PGR, and culture under photoperiodic conditions. Seven out of 30 trees tested showed embryogenic response but at lower frequencies (0.3% to 3.6%) than those obtained with adult *Q. suber* trees; the collection date also influenced the embryogenic response (Toribio et al. 2004; Valladares et al. 2006). The

scheme applied to SE of *Q. robur* differed from that used for *Q. suber* (Hernández et al. 2003a) in that the culture media were based on MS mineral components and a lower concentration of PGR in the induction media, whereas Schenk and Hildebrandt (1972) macronutrients and higher concentrations of both BA (10 μ M) and NAA (50 μ M) were successful for leaves from adult cork oak.

Once it was confirmed that genotype and selection of responsive explants collected at the optimal time were successful criteria for embryogenic induction in mature oak trees, other alternatives were proposed for searching for totipotent cells or tissues in recalcitrant species. Culture of the shoot apical meristems may provide a solution to problems related to recalcitrance, although the survival rates were normally very low (Bonga et al. 2010). Successful induction of somatic embryos in pedunculate oak, based on the use of shoot apex explants from material derived from adult trees has recently been reported (San-José et al. 2010). In the latter study, induction of embryogenic cultures was achieved in shoot tip explants (2 mm long, comprising the apical meristem and 3-4 pairs of leaf primordia) and leaf explants (the two apical- most expanding leaves) excised from axillary shoot cultures of three out of the five *Q. robur* trees over 100 years old that were evaluated. Within each responsive genotype, both leaf and shoot tip explants produced embryogenic responses, and embryogenic frequencies were relatively low (1.7%-12%) for both explant types, as occurred in the case of leaf explants excised from forced epicormic shoots of mature trees. Interesting, the failure to induce SE from shoot tips isolated from forced epicormic shoots of the corresponding trees suggest that the sterilization process has a negative effect on this type of explant. The use of shoot multiplication cultures to initiate the embryogenic process offers advantages over the use of explants from forced epicormic shoots from field-grown trees, as it ensures the availability of plant material all around the year and avoids the need to sterilize the plant tissues (San-José et al. 2010). The impact of genotype appears to be one of the limiting factors for the induction step of the SE process, as documented for other *Quercus* species (Hernández et al. 2003b; Vengadesan and Pijut 2009). Given that the frequencies of SE obtained in leaf explants from mature trees were not very different from those achieved using leaf explants from seedlings (Cuenca et al. 1999), the juvenile condition of the source tree, at least in leaf tissue, does not appear to be as crucial a factor as the genotype. A similar embryogenic response was observed between leaves from cork oak seedlings and from adult trees, as a result of the use of presumably reinvigorated or rejuvenated condition of the epicormic shoots (Hernández et al. 2003a).

With regard to the suitability of floral tissues for initiating SE, the study of Bueno et al. (1997), in which microspore-derived embryos of cork oak were obtained by combining a sucrose starvation treatment with a mild heat shock at 33° C for 5 days followed by culture at 25°C in a medium lacking PGR, is worthy of mention. After only 20 days of culture, embryos

were observed growing from the interior of the anthers, breaking free from the degenerating anther walls. These results indicate that stress may be the major signal for inhibition of gametophytic development of the microspores in favor of the embryogenic pathway. Ploidy level analysis, carried out on long-term (up to 12 months) microspore-derived embryo cultures, showed that most of the embryos (90.7%) were haploid, corresponding to their microspore origin, although a low percentage (7.3%) of diploid embryos was also observed (Bueno et al. 2003). These results confirm the accepted assumption that morphological changes observed in stressed microspores appear to be a prerequisite for transition to the de-differentiated state required for embryogenic induction (Bonga et al. 2010). Furthermore, a remarkable difference with respect to the induction of SE from leaf explants of the same species (Hernández et al. 2003a) must be highlighted, as no PGR were required to induce the androgenic process, and embryos were developed in shorter periods of time.

Although most of the studies carried out during the last decade refer to *Q. suber* and *Q. robur*, a few have involved the induction of somatic embryogenesis in holm oak (*Q. ilex*) (Mauri and Manzanera 2003) and red oak (*Q. rubra*) (Vengadesan and Pijut 2009). In both species, immature zygotic embryos were used as explants, giving rise to an induction frequency of 4.3% in the former, whereas a higher response (66%) of embryogenic callus was recorded in the latter. In addition, the induction of SE in *Q. alba* (white oak) following a procedure similar to that defined for pedunculate oak (San-José et al. 2010) was recently demonstrated (unpublished). Shoot tip explants and leaf explants excised from shoot cultures derived from selected 5-year-old trees produced somatic embryos at higher frequencies (up to 47%) than those reported for *Q. robur*. Interestingly, significant differences between explant types were observed, with a higher embryogenic response produced in the first expanding leaf from the apex.

There is evidence that soluble signal molecules may control cell differentiation in plant SE and that conditioned culture media from embryogenic cultures can promote SE. Signal molecules such as extracellular proteins (endochitinases), arabinogalactan proteins (AGPs) and lipochitooligosaccharides (LCOs) have been found to be important for the development of the embryogenic process (von Arnold et al. 2002). The effect of different concentrations of AGPs from both larch wood (Ben Amar et al. 2007) and gum arabic (Letarte et al. 2006) on initiation of somatic embryos in *Q. robur* and *Q. alba* was tested. Preliminary results indicate that the addition of these components to the culture media enhances the embryogenic response in both species.

Somatic embryo development in original explants When leaf explants or shoot apex explants derived from mature *Q. robur* trees were cultured in induction medium M1, they initially

responded by developing callus tissue at frequencies depending on the genotype. Callus proliferation was observed within two weeks and was mainly differentiated on the petiole stump region of leaf explants or cut surface of shoot tip explants, then progressing towards the distal ends. The SE followed an indirect pathway from this callus tissue formed in the original explants. In contrast, no callus formation was reported to occur in cork oak leaves where somatic embryos appeared almost directly at the leaf surface, without a defined pattern of arrangement (Hernández et al. 2003a). The timing of somatic embryo development was also influenced by genotype; some somatic embryos first began to appear towards the end of culture on M2 medium (9-10 weeks after culture initiation), although most emerged during culture in M3 expression medium, whereas in other genotypes, embryos began to become evident following transfer of the explants to M3 medium (12-15 weeks after culture initiation). Embryogenic cultures consisted of creamy-translucent nodular structures and somatic embryos, which developed from the callus tissue in both types of explant; induction of embryogenic tissues appears to require dissociation and partial necrosis of the specific callus regions. The embryogenic nodular structures observed in *Q. robur* were probably similar to the translucent nodules observed in other *Quercus* species, including *Q. ilex* (Féraud-Keller and Espagnac 1989) and *Q. suber* (El Mâataoui et al. 1990), or to the nodular proembryogenic masses described in other related species such as chestnut (Carraway and Merkle 1997; Corredoira et al. 2006a). In *Q. alba*, nodular embryogenic structures also originated from the callus tissue developed on the abaxial surface of leaf explants, and arose faster than those reported in pedunculate oak and cork oak, as the first embryogenic structures were evident during culture in M1 induction medium (6-8 weeks of culture initiation). Embryogenic nodular structures were embryogenic in character as they originated from embryogenic cell aggregates and in turn generated somatic embryos or other nodular structures as a form of repetitive embryogenesis.

Anatomical study Research on embryogenic cell initiation and early developmental stages of primary embryos induced in leaf explants (Corredoira et al. 2006b) and shoot tip explants (San-José et al. 2010) confirmed an indirect SE process. During culture in M1 induction medium, explants yielded callus tissue comprising parenchymatic cells with vascular elements, and actively dividing meristematic cells produced superficial layers of callus protuberances; these meristematic zones generally evolved into differentiated parenchyma cells rather than somatic embryos. At the end of culture in M2 medium and during culture in M3 medium, certain cells acquired an embryogenic character; these cells appeared interspersed with parenchyma cells, which were undergoing further vacuolization and degradation, in superficial callus regions. The formation of embryogenic cells was also associated with extensive division of perivascular cells in differentiating vascular bundles in the callus. In white oak, the formation of embryogenic

cells appears to be associated with a felodermis-like meristem or diffuse cambium established in superficial layers of the callus; in this case, cells exhibiting embryogenic-like cytological features also displayed intense accumulation of phenolic substances. Phenol accumulation in cells surrounding somatic embryos has been reported in myrtle (Canhoto et al. 1999) and may play an unknown role during early stages of SE in eucalyptus (Pinto et al. 2008). It is possible that this may account for the physical or physiological isolation of potential embryogenic cells as one of the pre-requisites for triggering the embryogenic program. Successive division of the embryogenic cells led to the formation of a few-celled proembryos and embryogenic aggregates within a thick common cell wall, indicating a probable unicellular origin, although a multicellular origin from bulging embryogenic areas cannot be discounted. Embryogenic aggregates gave rise to well-formed somatic embryos or to embryogenic nodular structures, which were formed from small vacuolated cells and meristematic areas, but lacked clear bipolarity. These cells were able to produce somatic embryos of multicellular origin from their meristematic areas, and somatic embryos of unicellular origin from embryogenic cells shed from the outermost layers. Somatic embryos, including cotyledonary stage embryos with shoot and root meristems, were apparent at successive stages of development.

Overall, these results suggest that leaf tissues are suitable explants for initiating SE in mature trees in different *Quercus* species, although pedunculate oak appears to be more recalcitrant than cork oak and white oak and the results obtained with the former closely resemble those obtained in leaf explants of a 50-year-old *Q. ilex* tree (Féraud-Keller and Espagnac 1989). Even in the case of shoot tip explants, the apical meristem itself generally senesced and died during culture, and the basal leaf primordia attached to the axial zone were the real source of generating calluses and, subsequently, embryogenic tissues. This suggests that the production of somatic embryos from oak shoot tip explants may be considered as a particular case of SE in which leaf primordia and their axillary basal meristems were involved. This is especially evident in *Q. alba*, where basal leaf primordia included in the shoot tip explants were the source of somatic embryos produced by this type of explant.

Secondary embryogenesis Once embryo initiation was achieved, even though induction frequencies were low, a large number of somatic embryos can be obtained by secondary embryogenesis (Toribio et al. 2004; Valladares et al. 2006). After isolation of somatic embryos from the original explant and culture in proliferation media, a recurrent embryogenesis mechanism is triggered, thus enabling the formation of secondary embryos giving rise to clonal embryogenic lines. Secondary embryos were mainly formed on the hypocotyl or root zone of the primary embryo, although some also appeared on the cotyledons. Production for scaling up and long-term maintenance of SE in different species of *Quercus* depends on their embryogenic

capacity through secondary embryogenesis. A semisolid proliferation medium supplemented with either a low concentration of cytokinins and auxins, or media without PGR were used for multiplication of embryogenic lines derived from pedunculate oak (Cuenca et al. 1999; Wilhelm 2000) and cork oak (Hernández et al. 2003b). For embryogenic lines derived from mature *Q. robur* trees, a proliferation medium consisting of MS basal medium including 500 mg/l casein hydrolysate, 0.44 μM BA and 0.27 μM NAA was initially adopted, and has made possible the continuous supply of somatic embryos during several years (Valladares et al. 2006; Sánchez et al. 2008). Further optimization experiments demonstrated that PGR treatment significantly affected proliferation and quality of the embryos (Mallón et al. 2011) with the best rates achieved when the proliferation medium was supplemented with 0.44 μM BA alone, resulting in significantly higher total numbers of embryos and of cotyledonary-stage embryos (used for germination purposes). Embryo production was affected by genotype, as expected given the impact of genotype on the induction step and germination step of the oak embryogenic systems. A genotypic effect was also documented in the proliferation of embryogenic lines of *Q. suber*, based on significant differences in the relative increase in fresh weight and number of ‘detachable’ embryos (Hernández et al. 2003b).

The histological origin and structural organization of the secondary embryogenesis process were investigated in *Q. robur* and *Q. suber*. A multicellular origin of embryos, and both direct and indirect secondary embryogenesis responses were reported in *Q. robur* (Zezgouti et al. 2001). In the case of the cork oak system (maintained in PGR-free medium), secondary embryos arose either in a multicellular budding pathway from a compact mass of proliferation or from isolated single cells (unicellular origin) in friable callus induced with relatively high levels of 2,4-D (Puigdejarrols et al. 1996, 2001).

Maturation, germination and plantlet conversion As in many other woody species, the low rate of conversion into plantlets is a major limitation for large-scale application of oak somatic embryogenesis. Maturation has been hampered in many species by repetitive embryogenesis, immaturity, embryo dormancy and precocious germination. During the maturation stage, somatic embryos should accumulate specific storage products at the appropriate time, as occurs in zygotic embryos (von Arnold et al. 2002). Absciscic acid (ABA) plays a fundamental role in processes such as regulation of the synthesis of storage proteins and late embryogenesis-abundant (LEA) proteins and water-stress-induced gene expression (Dodeman et al. 1997), in the reduction of secondary embryogenesis and in the inhibition of precocious germination (Kernode 1995). Desiccation, at least in zygotic embryos, is known to play a role in switching from the embryogenic to the germination phase. Reduction of intra- and intercellular-water

content by means of osmotic stress of the culture media may promote healthy development by mimicking the natural course of zygotic embryogenesis (Compton and Gray 1996).

Several approaches have been carried out to improve germination (root growth) and plantlet conversion (root and shoot growth) of somatic embryos in different oak species. Although the germination response, in terms of root development, may achieve values up to 80% in various oak species, many of the germinating embryos only exhibited root development, and the conversion rates were still not satisfactory (3% to 32%, Sánchez et al. 2003). A maturation stage allowing synthesis and accumulation of storage compounds and acquisition of desiccation tolerance probably also has to be completed by oak somatic embryos prior to germination. One form of inducing partial desiccation is to use an appropriate concentration and type of carbohydrate in the culture medium in order to induce osmotic stress. The culture of somatic embryos, prior to germination, in maturation media supplemented with 6% sorbitol or 6% mannitol plus 2-3% sucrose for one month, significantly favored the conversion rates in *Q. robur* embryogenic lines derived from leaf explants of seedlings (Sánchez et al. 2003), and in lines derived from mature trees (Valladares et al. 2006). These authors have also demonstrated that germination and conversion ability greatly depended on genotype; after 8 weeks of culture in germination medium, embryos of one genotype failed to germinate, and they continued to produce large numbers of secondary embryos while other genotypes exhibited conversion rates of up to 70% (Valladares et al. 2006). However, García-Martín et al. (2001) pointed out that the somatic embryo response of *Q. suber* was not improved by using a high concentration of sucrose prior to germination, unless accompanied by two months of chilling. Osmotic stress induced by increasing the concentration of agar (1%) in the maturation medium, with subsequent partial desiccation (moisture loss of 20-25%) promoted maturation and subsequent germination of pedunculate oak somatic embryos of culture lines of zygotic origin, thus achieving conversion rates up to 63% (Sunderlikova and Wilhelm 2002; Prewain et al. 2004). These authors also pointed out that a progressive increase in the concentration of proline and a 50% decrease in the concentration of ABA were observed to occur during embryo maturation (Prewain et al. 2004). There are no conclusive results on the exogenous application of ABA to promote maturation and subsequent germination of oak embryos. Although the addition of ABA to the culture media significantly reduced unwanted recurrent embryogenesis in *Q. ilex* (Mauri and Manzanera 2004) and significantly stimulated somatic embryo maturation in *Q. suber* (García-Martín et al. 2005), the treatments carried out in cork oak (Toribio et al. 2005) and in pedunculate oak did not have significant effects on germination and plantlet conversion.

For a number of woody species, culture of somatic embryos in a maturation medium must be followed by a period of cold storage to break embryo dormancy, a process associated with changes in the endogenous levels of ABA and gibberellic acid, (GA₃) (Deng and Cornu 1992).

The application of a 2-month cold period as a pre-germination treatment stimulated embryo germination and conversion in material from *Q. suber* of immature zygote origin (Fernández-Guijarro et al. 1995; García-Martín et al. 2001; Gonzalez-Benito et al. 2002a) and *Q. ilex* (Mauri and Manzanera 2004), as well as in embryogenic lines from *Q. suber* of mature tree origin (Hernández et al. 2003b). Similarly, the application of activated charcoal and a mixture of glutamine, arginine and asparagine coupled with cold storage at 4°C for two months had a favorable effect on plantlet conversion in cork oak (Pintos et al. 2010). In the embryogenic system derived from immature cotyledons of *Q. rubra*, germination was performed by desiccation for 3 days followed by cold storage for 8 weeks (Vengadesan and Pijut 2009). However, two months of storage at 4°C between maturation and germination treatments did not have any significant effect on the conversion frequency in pedunculate oak embryos from mature trees, although germination, in terms of root-only development, was promoted in the three embryogenic lines investigated (Martínez et al. 2008). Overall, stratification appears to be an important treatment for promoting germination in oak somatic embryos, as occurred in chestnut, another recalcitrant seed species and member of the family *Fagaceae* (Corredoira et al. 2003; Andrade and Merkle 2005).

Mature somatic embryos of *Q. robur* were usually germinated in media supplemented with 0.28-0.56 μM BA with or without IBA (0.1 μM) (Cuenca et al. 1999; Sunderlikova and Wilhelm 2002; Valladares et al. 2006), whereas germination media devoid of PGR were used with *Q. suber* (Hernández et al. 2003b) and *Q. ilex* (Mauri and Manzanera 2004). A positive effect of inclusion of GA_3 in the germination medium was reported for *Q. acutissima* somatic embryos (Kim 2000), which is consistent with the results obtained by Vengadesan and Pijut (2009) in *Q. rubra* somatic embryos, in which the combination of BA (0.44 μM) and GA_3 (0.29 μM) favored simultaneous development of both the shoot and the root, thus yielding a high frequency of germination (61%). However, preculture of *Q. robur* somatic embryos in GA_3 (8.65 μM) medium for two weeks prior to transfer to germination medium did not improve the conversion rates, but did reduce root length and the number of leaves per regenerated plantlet (Sánchez et al. 2003).

Epicotyl and shoot development in oak germinating embryos appears to be a limiting factor, and plantlet conversion is still not operational on a large scale. The shoot promoting ability of thidiazuron (TDZ) suggests that this may influence the organization and development of what would otherwise be a poorly developed shoot apical meristem, and the effect of this compound was therefore investigated during the germination step in *Q. robur* (Martínez et al. 2008). Addition of 0.05-0.1 μM TDZ to the germination medium for an initial period of seven days induced multiple shoot formation in the epicotyl region (the main shoot plus cotyledonary axillary shoots) of the germinating embryos, increasing the plantlet conversion frequency and

the shoot-only elongation frequency, especially in the case of hard-to-convert embryogenic lines. Shoots excised from shoot-only germinating embryos can be elongated, multiplied and rooted by proven micropropagation procedures via axillary shoot proliferation (Martínez et al. 2008), as already described in this review. This procedure would also provide an interesting tool to study if rejuvenation of the regenerated shoots through somatic embryogenesis (induced from mature tissues) occurs. The combined strategy based on somatic embryo conversion to the extent that it is possible, and proliferation and rooting the shoots of unconverted embryos that produce them, has been proposed for production of plants from embryogenic transgenic lines of European chestnut and American chestnut (Maynard et al. 2008). The initial experiments involving application of TDZ to germinating embryos also gave rise to an additional finding that adventitious bud regeneration was induced on the cotyledons of TDZ-treated embryos (Martínez et al. 2008), which may be useful in genetic transformation procedures applied to this species.

In an attempt to understand the maturation and germination processes in pedunculate oak, the expression patterns of the storage protein legumin gene and Em- and dehydrin-like homologues were investigated (Sunderlikova and Wilhelm 2002; Sunderlikova et al 2009a). The latter authors concluded that these proteins may be useful as markers of the physiological and maturation condition of somatic embryos (see also Genomics section).

Data from field experiments with plants derived from somatic embryos are scarce and limited to *Q. suber* plants regenerated from mature selected trees, from their half-sib progenies by somatic embryogenesis and from zygotic seedlings of the same families. Differences as a result of genotype and family effects were notable and no apparent morphological alterations were detected among the trees of tested progenies (Celestino et al. 2007).

Genetic stability

The genetic stability of in vitro regenerated plants is an essential requisite for large-scale clonal forestry. Evaluation of genetic fidelity in plantlets of the genus *Quercus* obtained through somatic embryogenesis is of vital importance because in these species, as a result of their long life cycle, assessment and detection of possible somaclonal variation is critical to prevent the spread of instability among high-value genotypes. PCR-based markers (RAPDs, AFLPs, SSRs) and cytological studies have been used to evaluate the potential genetic variation in embryogenic cultures and regenerated plants of different *Quercus* species.

Four embryogenic lines initiated from explants of 4-week-old *Quercus robur* seedlings maintained for more than 4 years by secondary embryogenesis were subjected to RAPD analysis. In addition to these lines, a line derived from a nodular embryogenic callus, which ceased to be embryogenic after a few subculture cycles, was also analyzed (Sánchez et al.

2003). No polymorphism was detected within and between embryogenic lines originated from different types of explant from the same seedling, but enough polymorphism was detected among genotypes originating from different plants to enable the origins to be distinguished. There were also no differences in DNA sequences between regenerated plantlets and their somatic embryos of origin. However, these authors observed genetic variation between the callus line that had lost its embryogenic capacity and the three embryogenic lines derived from the same seedling (Sánchez et al. 2003). The genomic alteration appears to have caused loss of the capacity for embryogenesis. Callus proliferation probably involves fewer active genes than the formation of somatic embryos or converted plantlets, and mutations in these active genes may be the cause of the loss of ability to undergo secondary embryogenesis (Heinze and Schmidt, 1994). De Verno et al. (1994) suggested that the embryogenic tissue may be composed of a mixture of altered and unaltered cells and genomes, and that only the unaltered genome would be able to regenerate plantlets. In addition, five microsatellite loci were used to test the genetic stability of three somatic embryogenic lines derived from immature zygotic embryos of *Q. robur* and derived plantlets (Wilhelm et al. 2005). Variation in DNA was detected among somatic embryos within the three embryogenic lines with a high mutation frequency of 29.2–62.5%, depending on the genotype, whereas no genetic instability was found among the regenerated plantlets. The authors, in line with the results reported by Sánchez et al. (2003), suggested that regeneration would be a selective process in which conversion into plants would be prevented in somatic embryos with mutations. Both RAPD and AFLP markers were used to record differences between embryogenic lines, even between those that arose from half-sib zygotic embryos, but no genetic variation was found among somatic embryos within embryogenic lines of zygotic embryo origin of *Q. suber* (Gallego et al. 1997; Hornero et al. 2001).

The genetic stability of in vitro cultures and regenerated plants compared with the original selected material from which they are derived is essential for the maintenance and propagation of elite genotypes. RAPD analysis was used to check the genetic stability of somatic embryos induced from mature *Q. robur* tissues. No evidence of genetic changes was found within or between those embryogenic lines and the mature trees of origin, and stability was also maintained between the regenerated plantlets and trees of origin (Valladares et al., 2006). However, pre-existing variation among tissues from the donor plant was attributed to DNA instability detected among AFLPs in *Q. suber* somatic embryos induced from leaves of adult trees (Hornero et al. 2001). Similarly, Lopes et al. (2006) used microsatellites to demonstrate mutations in *Q. suber* somatic embryos derived from leaves from a mature field tree. Nevertheless these authors found no somaclonal variation in another embryogenic line of adult origin.

Therefore, to certify the genetic stability of regenerated plants by in vitro culture, it is important to use a variety of approaches such as flow cytometry, since PCR-based markers fail to detect variation at the ploidy level. Endemann et al. (2001) used flow cytometry analysis to detect tetraploidy in 8% of pedunculate oak somatic embryos originated from zygotic embryos and maintained by secondary embryogenesis, and the changes in chromosome number appeared to be correlated with the duration of the in vitro culture. Using the same technique, Loureiro et al. (2005) found no polyploidy or significant variations in DNA content among cork oak somatic embryos, and between the embryos and the mature mother trees. These authors also observed the same ploidy level when the somatic embryos were germinated and converted to plantlets.

Pre-existing variation in the original explants, probably linked to the genotype, and de novo induced mutational events during the embryogenic process, appear to be the most important aspects related to the genetic stability of oak somatic embryos (Wilhelm et al. 2005).

Cryopreservation

Gene conservation programmes are recommended for oak species because the genetic resources are threatened by multiple factors such as the introduction of exotic genotypes, species purification, neglected practices, diseases, etc. In situ conservation methods are generally preferred, although if natural methods are not sufficient, ex situ conservation through clonal seed orchards should be used (Ducousso and Bordacs 2004). Furthermore, as pointed out in the Introduction section, *Quercus* species are late-maturing and late-flowering, exhibit irregular seed set, and produce seeds that are recalcitrant to conventional conservation procedures. Recalcitrant seeds have a high water content and can be only stored for relatively short periods of time because they are sensitive to desiccation.

There are several potential advantages of using synthetic seeds, including the potential for long-term storage and for automation and commercialization purposes, but there is scant information on the feasibility of encapsulation in *Quercus* species. In one study, apical buds of *Q. robur* and *Q. cerris* were encapsulated in an alginate matrix, giving rise to successful regeneration after 6 weeks of cold storage at 4°C (Tsvetkov and Hausman 2005). Furthermore, the production of synthetic seeds and their ability to be stored has been investigated in somatic embryos of *Q. robur* (Prewein and Wilhelm 2003) and *Q. suber* (Pintos et al. 2008). However, the results obtained in both studies showed that conversion rates were considerably reduced after 2-3-months of cold storage at 4°C, and only short-term storage of encapsulated embryos was possible.

In general, strategies for forest biodiversity conservation are today well-defined (Häggman et al. 2008), among which cryopreservation is viewed as a complementary, secondary storage method designed as a secure backup for living collections (Reed 2008). Well-defined in vitro techniques are required to recover the multiplication capacity of supercooled material via axillary shoot growth or somatic embryogenesis. Only under these conditions, the application of cryopreservation in the development of large gene banks is justified. Cryopreservation is currently routinely applied to embryogenic lines awaiting field testing results (Sutton 2002) and the first approaches on the application of cryopreservation to hardwood forest tree species have recently been published (Harvenge et al. 2004; Vidal et al. 2010a).

Zygotic embryo axes of pedunculate oak have been cryopreserved using the simplest desiccation-based method, by means of fat flash drying and direct immersion in liquid nitrogen (LN), which yielded appreciable germination recovery rates (Berjak et al. 2000). These rates were not improved when more complex dehydration procedures were applied. Better results were reported with embryonic axes of *Quercus faginea*, a tree that occurs widely in the Iberian peninsula (González-Benito and Pérez-Ruiz 1992). Prior to storage in LN, the axes were desiccated either under sterile conditions in a laminar air flow bench for different periods of time, or by soaking them in 15% dimethylsulfoxide (DMSO) for 1 h and subsequently immersing them in LN. The best recovery rates (60% germination) were obtained after desiccation for 3 h, after which the moisture content decreased from the initial value of 63.7% to 12%. However, attempts to subject cork oak zygotic embryo axes to cryostorage by partial desiccation followed by rapid immersion in LN were unsuccessful, as only unorganized post-cooled growth of axes was reported (González-Benito et al. 2002b).

As progress in the induction of somatic embryogenesis from selected trees has been evident in the last years in species such as pedunculate and cork oak, interest in the cryopreservation of embryogenic lines has increased. Cryopreservation techniques referred to as ‘vitrification-based techniques’ are now applied to different oak species. Briefly, embryo clumps (4-6 mg) of globular or heart-shaped secondary embryos isolated from embryogenic lines were pre-cultured for 3 days on a solid MS medium containing 0.3 M sucrose, and were then transferred to fresh medium containing 0.7 M sucrose for another 4 days. The embryos were placed in cryovials, then subjected to vitrification by application of PVS2 solution (30% w/v glycerol, 15% w/v DMSO and 15% w/v ethylenglycol in MS medium containing 0.4 M sucrose) for 30 min at 25°C, resuspension in the PVS2 solution and then plunging in LN. After rewarming, recovery rates (relative to the generation of new embryos from surviving cells of the cryopreserved clumps by repetitive embryogenesis) of 75%-80% were recorded for somatic embryos of pedunculate oak (Martínez et al. 2003; Sánchez et al. 2008). Following a similar procedure, higher recovery rates (>90%) were achieved with different embryogenic lines initiated from

mature selected trees of cork oak (Valladares et al. 2004). Preculture on a medium containing sugars appeared to be an essential step for successful cryostorage in the vitrification procedure. A high concentration of sucrose may increase tolerance to desiccation, which may stimulate the accumulation of certain sugars in the cells. The effect of increased sugar content on cell integrity may cause a decrease in cell volume as a result of osmotic process and direct stabilization of membranes (Percy et al. 2001).

Construction of a large gene bank of cork oak genotypes by use of the cryopreservation methodology developed at laboratory level is a reality. Currently, 100% of the 51 embryogenic lines corresponding to genotypes of both juvenile (44 genotypes) and mature (7 genotypes) origins incorporated to the gene bank recovered their capacity to proliferate after immersion in LN. Furthermore, three experimental plots have been established with the selected materials to determine their productivity and heritable characters by investigation of progenies. During field testing, copies of the material will be preserved under LN (Vidal et al. 2010a).

One concern related to cryopreservation methods is the genetic stability of the post-cooled and regenerated plants. The potential somaclonal variation of in vitro tissue cultures, as well as the mutagenic potential of the most popular cryoprotectant component of the PVS2 vitrification solution, DMSO, are well-known, although to date there is no evidence of morphological or genetic alterations in forest trees as a result of cryopreservation (Häggman et al. 2008). Aronen et al. (1999) reported genetic changes in DMSO-treated, non-supercooled samples, but not in DMSO-treated and cryostored samples of *Abies cephalonica* somatic embryos during cryopreservation. The authors speculated that, although cryostorage does not remove the mutagenic potential of DMSO, it probably eliminates a high proportion of the cells bearing genetic changes. In our experience, RAPD profiles of 5 embryogenic lines of *Quercus robur* cryopreserved for years, and the corresponding regenerated plantlets were identical to those of the controls and no genetic instability (sporadic mutations or genetic reordering) was detected (Sánchez et al. 2008).

Genetic transformation

As pointed out above, the induction and proliferation of somatic embryos derived from both juvenile and mature material of different *Quercus* species has been demonstrated. Factors affecting the germination and conversion of somatic embryos into plantlets were also studied and, consequently, the most appropriate target material for the application of gene transfer is now available.

Genetic transformation protocols for cork oak embryogenic lines induced from both mature (Alvarez et al. 2004) and juvenile (Sánchez et al. 2005) materials through co-culture with different strains of *Agrobacterium tumefaciens* harbouring plasmids containing marker genes

have been defined. An optimized protocol for *Agrobacterium*-mediated transformation of mature embryogenic masses of this species, in which variables such as the genotype, explant type and the preculture period were studied, has also been reported (Alvarez and Ordás 2007). Genotype has a strong influence on the transformation frequency, as only 3 out of the 6 embryogenic lines tested were kanamycin-resistant after 4 months of culture in selective media. The transformation efficiencies were highly variable among experiments, which may be related to the physiological state of the somatic embryo clumps used in the experiments. Unfortunately, the plantlet conversion rates were not reported in any of these studies. In addition, cork oak embryogenic masses were transformed using *Agrobacterium* strain AGL1, which carried the plasmid pBINUbiBar, harbouring the genes *nptII* and *bar*, the latter under control of the maize ubiquitin promoter. Varying activities of phosphinothricin acetyl transferase were detected among the lines, which carried 1-4 copies of the insert. The stability and expression of the transgenes 3 months after thawing of the cryopreserved transgenic embryogenic lines were confirmed by molecular and biochemical probes. Herbicide resistance has only been tested at laboratory level, as plantlet conversion has not been reported (Alvarez et al 2009).

Initial attempts to obtain transformed plants of pedunculate oak (*Q. robur*) were unsuccessful (Roest et al. 1991; Wilhelm et al. 1996). A protocol for genetic transformation of somatic embryos derived from juvenile (2 lines) and mature (3 lines) pedunculate oak trees co-cultured with the *Agrobacterium* strain EHA105 harbouring a p35SGUSINT plasmid containing a *nptII* marker gene and a *uidA* reporter gene, has recently been reported (Vidal et al. 2010b). Histochemical and molecular analyses confirmed the presence of *nptII* and *uidA* genes in the roots and leaves of transformed plants. Transformation efficiencies ranged from up to 2%, in an embryogenic line derived from a 300-year-old tree, to 6% in a juvenile genotype. Twelve independent transgenic lines were obtained and transgenic plantlets were recovered and are growing in a greenhouse.

The feasibility of genetic transformation of pedunculate oak and cork oak somatic embryos by means of the co-culture technique with several *Agrobacterium* strains has been demonstrated. Taking into consideration these advances, the next step should be to study the response of these or other oak species after genetic transformation with genes of interest, as carried out with chestnut (Maynard et al. 2008), a member of the family *Fagaceae*.

Genomics

Genomic information on many forestry species has increased dramatically in recent years and genomic resources available for tree improvement and domestication programs include genomic sequencing, EST sequencing, transcript profiling, gene transfer, molecular markers for mapping and breeding, etc. (Merkle and Nairn 2005). With advances in molecular biology and

identification of adaptive genes, differences in gene sequences for genes involved in the control of a particular adaptive trait would be used for assessing diversity and adaptiveness (Savill et al. 2005). Current progress in tree genomics is so rapid that few studies have been published to date and much documentation is often limited to electronic resources. There are a number of websites related to genomics and gene discovery in trees (Nehra et al. 2005). An integrated web-based resource for members of the *Fagaceae* family, including *Quercus* (Fagaceae Genomic Database (FGD) <http://www.fagaceae.org>), has been developed. This site disseminates data and analyses by providing relevant sequence information, homology results, genetic/physical map information, SSRs, SNPs; and other genomic data are being posted as they become available. The most important information from studies involving the genomics of *Quercus* species, including construction of genetic linkage maps, microsatellite markers, EST-derived markers, transcriptome investigations and differential gene expression for the identification of genes involved in the adaptation of oak trees and QTL detection for phenotypic traits that respond to natural selection and to specific stresses, has been reviewed in detail (Kremer et al. 2007).

Most research activities involving the molecular genetics of *Quercus* have addressed population genetics, unlike in other forest species, for which tree improvement has been the main goal. Two oak mapping initiatives have been carried out: for European white oaks the objective is to detect QTLs that control morphological and adaptive traits involved in species differentiation (Petit et al. 2004; Saintagne et al. 2004; Scotti-Saintagne et al. 2004a), while in American red oaks, the ultimate goal is the detection of QTLs and genes that control heartwood color and resistance to specific pests (Aldrich et al. 2002).

Implementation studies have been carried out on transcriptomic investigations and differential gene expression to identify genes involved in the adaptation of oak species to their environments. Genetic variation of bud burst and early growth has been estimated in a full-sib family of *Q. robur* comprising 278 offsprings. Bud burst appears to be controlled by several QTLs with rather low to moderate effects and height growth depends on fewer QTLs with moderate to strong effects (Scotti-Saintagne et al. 2004b). After this, 801 ESTs were generated derived from 6 developmental stages of bud burst. Microarray and real-time RT-PCR showed that several differential expressions between quiescent and swelling buds constitute relevant candidates for the signaling pathway of bud burst in trees (Deroy et al. 2006).

Forest trees are threatened by changing environmental conditions because global climate change. Thirteen osmotic stress-induced genes of *Quercus* ssp. were included in genetic linkage maps of oak (Porth et al. 2005) and later studies on QTLs of tolerance to waterlogging in pedunculate oak identified significant clusters of QTLs, at inter- and intra-trait level, suggesting the occurrence of a genetically controlled cascade response to waterlogging (Parelle et al. 2007).

Quantitative trait loci controlling water use efficiency and related traits were also studied in *Q. robur* (Brendel et al. 2008).

Vegetative propagation using either conventional cutting procedures and micropropagation techniques, are used for the deployment of improved genetic forest tree species. Assessment of rooting in cuttings of pedunculate oak allowed the detection of 10 QTLs explaining between 4% and 13.8% of the phenotypic variance, of which the strongest were stable across years (Scotti-Saintagne et al. 2005). Furthermore, a full-length cDNA clone named QrCPE, which is differentially expressed during in vitro culture of mature (difficult-to-root) and juvenile (easy-to-root) shoots of pedunculate oak was identified by differential display (Gil et al. 2003). The deduced amino acid sequence showed that QrCPE encodes a small putative cell wall protein that is rich in glycine and histidine residues. Accumulation of QrCPE mRNA was higher in mature than in juvenile microshoots at the end of the multiplication and rooting steps. QrCPE accumulates in ontogenetically older organs of oak trees, although it is present in zygotic and somatic embryos but absent from callus cells. Expression of QrCPE gene was also investigated in juvenile-like microshoots during the in vitro rooting process, after demonstration of the auxin-mediated induction of this gene in rooting competent shoots (Covelo et al. 2009). On the other hand, an *Em*-like gene, QrEm, has been isolated from *Q. robur* zygotic embryos, which is expressed in the mid-maturation phase. Abundant transcript accumulation was observed when somatic embryos of the species were subjected to 6% sorbitol or after partial desiccation. The results of in situ hybridization experiments appear to suggest that the QrEm gene is embryo-specific and its expression is developmentally regulated (Sunderlikova et al. 2009a). Furthermore, three dehydrin genes, *QrDhn1*, *QrDhn2*, *QrDhn3*, were isolated from pedunculate oak. The *QrDhn1* gene was expressed during later stages of zygotic embryo development but only in somatic embryos when exposed to osmotic or desiccation stress. The other two dehydrin genes were expressed only in somatic embryos (both untreated and osmotically stressed), and the authors suggest that different dehydrins were involved in the process of seed maturation and response to altered osmotic (water status) conditions in somatic embryos (Sunderlikova et al. 2009b). These and other members of the dehydrin gene family may contribute to clarifying non-orthodox (oak) seed behavior and to identifying mechanisms related to desiccation tolerance. Finally, 17 differentially proteins have been identified in a proteomic study of *Q. suber* somatic and gametic in vitro culture-derived embryos. Some of the proteins identified are involved in stress and pollen development and others are associated with metabolism of tannins and phenylpropanoids (Gómez et al. 2009). Approaches using functional genomics to examine the molecular and cellular mechanisms that control organogenesis and somatic embryogenesis will contribute to better practical application of these processes to recalcitrant species such as oaks.

From the development of genetic maps in species of *Quercus*, *Castanea* and *Fagus* (all genera belonging to the family *Fagaceae*) carried out to date, it appears obvious that molecular markers could be easily transferred from *Quercus* to *Castanea* (Barreneche et al. 2004; Casasoli et al. 2006), but less easily to *Fagus*, and ongoing activities in comparative mapping suggest that there is a strong macrosyntheny between the phylogenetically close genera *Quercus* and *Castanea* (Kremer et al. 2007).

Conclusions

Conventional long-term tree improvement programs have not been carried out in the long-rotation species of the genus *Quercus*. The rapid development of biotechnological techniques, including in vitro tissue culture, gene transfer, conservation, molecular genetics associated with physiological and taxonomic studies, gene mapping, and identification of genes and alleles linked to specific traits, provides an opportunity for breeding, propagation, deployment and management applied to *Quercus* species. Although most of these species are classified as recalcitrant to regeneration through in vitro tissue culture, enormous progress has been made in the application of biotechnological tools to the most important species of *Quercus* during the last decade. Procedures of micropropagation through organogenesis and somatic embryogenesis using explants isolated from selected adult trees, and protocols for genetic transformation and cryopreservation of elite genotypes have been defined, allowing the regeneration of whole plants, regardless of the method used. However, the organogenic/embryogenic response is clearly dependent on the genotype and, in the embryogenic process, the low conversion rates of the somatic embryos into plantlets of most of the species/genotypes studied may be considered as one of the most important obstacles to the large scale propagation of selected oak genotypes. Most research involving genomics of *Quercus* species have concerned population genetics. Approaches using functional genomics to examine the molecular and cellular mechanisms that control organogenesis and or somatic embryogenesis are still scarce, and much effort should be made in this respect in the near future, to enable better application of oak clonal forestry. In addition, efforts should involve the isolation and characterization of genes related to other specific traits (i.e. disease and or pest resistance, wood quality, tree form), which would also be of interest for improving the species.

Acknowledgements This research was partly funded by the Xunta de Galicia (Spain) through project 09MRU002400PR.

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Table 1. Studies on micropropagation through axillary shoot proliferation in oak species of economic relevance

Species	Explant source/origin	Medium	Growth response	Reference
<i>Q. robur</i>				
	Embryonic axes and shoot tips and nodal explants from seedlings and stump sprouts	GD, BA (4.44-0.44) ¹ Dipping IBA (1 g/L) 2 min	Shoot proliferation Rooting	Vieitez et al., 1985
	Shoot tips from seedlings and grafts from mature trees	1/2MS, BA (0.44) IBA (4.9) 8 d	Shoot proliferation Rooting	Favre and Juncker 1987; Juncker and Favre 1989
	Shoot cultures from seedlings and stump sprouts	WPM, BA (2.66-0.88) or TDZ (0.0045-0.0090)	Shoot proliferation Rooting, acclimatization	Chalupa 1988
	Shoot cultures from seedlings and epicormic shoots from mature trees	GD, BA (0.88)	Shoot proliferation; recycling horizontal shoots	San-José et al. 1988
	Shoot tips, nodal segments from seedlings and stump sprouts	GD, BA (0.08) or Z (0.46) WPM, BA (2.66-0.88) WPM, IBA (0.98-4.90)	Shoot elongation, rooting Shoot proliferation Rooting, acclimatization	Chalupa 1993
	Shoot tips from forced epicormic shoots from mature trees	WPM, BA (4.4-2.2), AC (0.5%) IBA (4.6)	Shoot proliferation Rooting	Evers et al. 1993
	Zygotic embryos	Commercial rooting powder	Ex vitro rooting	Meier-Dinkel et al. 1993
	Shoot tips and nodal segments from forced epicormic shoots from mature trees	GD, BA (0.88)	Shoot proliferation; recycling horizontal shoots	Vieitez et al. 1994
	Shoot cultures from forced epicormic shoots from mature trees	IBA (14.8) 7 d 1/3GD, IBA (122.5) 24 h, AC (1%)	Rooting Rooting	Sánchez et al. 1996
	Shoot cultures from seedlings	1/2GD, IBA (4.90) 7 d	Rooting	Puddephat et al. 1999
<i>Q. petraea</i>				
	Shoot tips and nodal segments from seedlings and stump sprouts	GD BA (0.88) 1/2GD, dipping IBA (0.5 g/L) 6 min	Shoot proliferation Rooting	San-José et al. 1990
	Shoot tips and nodal segments from seedlings and stump sprouts	WPM, BA (2.66-0.88) WPM, IBA (0.98-4.90)	Shoot proliferation Rooting	Chalupa 1993
<i>Q. rubra</i>				
	Shoot tip and nodal	WPM, BA (4.44-0.88)	Shoot proliferation;	Vieitez et al. 1993

	segments from seedlings and from 4-y-old trees. Epicormic shoots from mature trees		recycling horizontal shoots	
	Shoot cultures	1/2WPM, IBA (122.5) 24 h, AC (1%)	Rooting	Sánchez et al. 1996
	Cotyledonary nodes from 8-w-old seedlings	WPM, BA (4.4), GA ₃ (0.29)	Shoot proliferation	Vengadesan and Pijut 2007
	Shoot tips and nodal segments from epicormic shoots from 7-y-old trees	WPM, IBA (4.9)	Rooting	
		WPM, BA (0.88-0.44), AgNO ₃ (3 mg/L)	Shoot proliferation, horizontal explants	Vieitez et al. 2009
		1/2WPM, IBA (122.5) 48 h, AC (0.4%)	Rooting	
<i>Q. suber</i>	Apical and axillary buds, flushed shoots from stem cuttings of mature trees	GD, WPM, BA (4.44-2.22), IAA (0.57)	Shoot proliferation	Romano et al, 1992
		Dipping IBA (0.5 g/L) 10 s	Rooting	
<i>Q. alba</i>	Shoot tips and nodal segments from epicormic shoots from 7-y-old trees	GD, BA (0.88-0.44), Z (2.28)	Shoot proliferation, horizontal explants	Vieitez et al. 2009
		1/2GD, IBA (122.5) 48 h, AC (0.4%)	Rooting	
<i>Q. bicolor</i>	Shoot tips and nodal segments from epicormic shoots from 7-y-old trees	GD, BA (0.88-0.44), Z (2.28)	Shoot proliferation, horizontal explants	Vieitez et al. 2009
		1/2GD, IBA (122.5) 48 h, AC (0.4%)	Rooting	

¹Quantities in brackets are expressed in µM unless otherwise stated

Mineral media: GD: Gresshoff and Doy (1972); MS: Murashige and Skoog (1962); WPM: Woody Plant Medium (Lloyd and McCown 1980)

Supplements: AC: Activated charcoal; BA: 6-Benzylaminopurine; GA₃: Gibberellic acid; IAA: Indole-3-acetic acid; IBA: 3-Indolebutyric acid; TDZ: Thidiazuron; Z: Zeatin

Table 2. Summary of somatic embryogenesis studies of *Quercus* species, with special reference to those published from the year 2000.

Species	Explant	Origin	Growth response/ Remarks	Reference
<i>Q. robur</i>	Leaves, internodes	S	Somatic embryos, plantlet development	Cuenca et al. 1999
	Leaf segments	MT	Somatic embryos, plantlet development	Chalupa 2000
	Embryogenic lines	ZE	Secondary somatic embryos, anatomical study	Zegzouti et al. 2001
	Embryogenic lines	ZE	Maturation, germination, plantlet development	Sunderlikova and Wilhelm 2002; Prewen et al. 2004
	Embryogenic lines	S	Maturation, germination, plantlet development	Sánchez et al. 2003
	Leaves	MT	Somatic embryos, plantlet development	Toribio et al. 2004
	Leaves	MT	Somatic embryos, plantlet development	Valladares et al. 2006
	Leaves	MT	Somatic embryo origin, anatomical study	Corredoira et al. 2006a,b
	Embryogenic lines	MT	Secondary somatic embryos, plantlet development	Sánchez et al. 2008
	Embryogenic lines	MT	Maturation, germination, plantlet development	Martínez et al. 2008
	Shoot tips and leaves	Shoot cultures from MT	Somatic embryos, anatomical study	San-José et al. 2010
	Embryogenic lines	MT	Secondary somatic embryos	Mallón et al. (in press)
<i>Q. suber</i>	Leaves	MT	Somatic embryos, plantlet development	Hernández et al. 2001
	Embryogenic lines	S	Maturation, germination	García-Martín et al. 2001
	Embryogenic lines	S	Secondary somatic embryos, anatomical study	Puigdejarrols et al., 1996, 2001
	Leaves	MT	Somatic embryos	Pinto et al., 2002
	Embryogenic lines	ZE	Germination	González-Benito et al. 2002a
	Leaves	MT	Somatic embryos	Hernández et al. 2003a
	Leaves	MT	Secondary somatic embryos, germination, plantlet development	Hernández et al. 2003b
	Anthers	MT	Microspore-derived embryos	Bueno et al. 2003
	Embryogenic line	S	Somatic embryo maturation	García-Martín et al. 2005
	Leaves	MT	Somatic embryos, maturation, germination, plantlet development	Toribio et al. 2005
<i>Q. rubra</i>	Somatic seedlings	MT	Acclimatization, field tests	Celestino et al. 2007
	Embryogenic lines	MT	Maturation, plantlet development	Pintos et al. 2010
<i>Q. ilex</i>	Immature cotyledons	ZE	Somatic embryos, maturation, plantlet development	Vengadesan and Pijut 2009
	Immature zygotic embryos	ZE	Somatic embryos, secondary somatic embryos, maturation	Mauri and Manzanera 2003

<i>Q. accutissima</i>	Embryogenic lines	ZE	Maturation, germination	Mauri and Manzanera 2004
	Immature, mature zygotic embryos	ZE	Somatic embryos, maturation, germination	Kim 2000

* S, seedlings; MT, mature trees; ZE, zygotic embryos