MINI-REVIEW

Hairy root type plant in vitro systems as sources of bioactive substances

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Abstract "Hairy root" systems, obtained by transforming plant tissues with the "natural genetic engineer" Agrobacterium rhizogenes, have been known for more than three decades. To date, hairy root cultures have been obtained from more than 100 plant species, including several endangered medicinal plants, affording opportunities to produce important phytochemicals and proteins in eco-friendly conditions. Diverse strategies can be applied to improve the yields of desired metabolites and to produce recombinant proteins. Furthermore, recent advances in bioreactor design and construction allow hairy root-based technologies to be scaled up while maintaining their biosynthetic potential. This review highlights recent progress in the field and outlines future prospects for exploiting the potential utility of hairy root cultures as "chemical factories" for producing bioactive substances.

Keywords Agrobacterium rhizogenes · Hairy roots · Secondary metabolites · Bioreactors

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Introduction

From time immemorial, mankind has been highly dependent on plants as sources of proteins, carbohydrates and fats (Rao and Ravishankar 2002). Furthermore, the volume and range of phytochemicals used by modern society (inter alia as drugs, nutrients, cosmetic additives and biopesticides) are continuously expanding. Currently, more than 25% of all prescribed medicines used in industrialized countries are derived either directly or indirectly from plants, and annual sales of these products in the USA alone exceeded 30 billion dollars in 2002 (Raskin et al. 2002; Fowler 2006). These high demands are driving efforts to develop new ways to produce plant-derived metabolites.

Plant in vitro techniques, in which plant cells, tissue and organs are cultivated in sterile conditions totally independently of geographical and climatic factors, offer such alternatives for producing important metabolites, and several technologies based on them have been developed (Kieran 2001). The exploitation of transformed root cultures (also called "hairy roots," by analogy with the natural plant disease of the same name, which causes massive production of adventitious roots with numerous root hairs) presents a relatively novel approach to in vitro plant biotechnology that has received increasing attention in recent years. Although the mechanism of Agrobacteriummediated genetic transformation has been known for more than 30 years, the enormous capacity of transformed root cultures has been largely neglected for most of that time. However, in the mid-1980s, hints of the biosynthetic potential of hairy roots were obtained through a series of investigations, focusing mainly on their alkaloids production (Mano et al. 1986; Payne et al. 1987). During the 1990s, research in this field entered an "exponential phase," and more than 400 papers on hairy roots were published

during this decade (source: SCOPUS database). The published studies highlighted several advantages of transformed root cultures, including their relatively fast growth rates (in hormone-free media), genetic and biochemical stability and capacity for organogenesis-associated synthesis of metabolites (Shanks and Morgan 1999; Sevon and Oksman-Caldentey 2002). In addition, recent developments in bioreactor systems provide ways in which it may be possible to scale up hairy root cultivation from small-scale systems to large-scale industrial processes (Guillon et al. 2006a). Furthermore, growing interest in hairy root systems from private companies (e.g. ROOTec bioactives GmbH, Basel, Switzerland), should facilitate the transfer of knowledge from academic laboratories to the pharmaceutical, food and cosmetic industries.

The present review summarizes recent developments in hairy roots research, including methods for mother plant transformations, the production of valuable biological active substances and strategies for enhancing the yields of desired metabolites. It also considers recent advances in bioreactor designs for hairy root cultivation and scaling-up processes.

Historical background and establishment

At the beginning of the last century, two plant diseases, crown-gall and hairy root, caused significant losses in vineyards, plum, apple and peach nurseries, and several investigations were initiated to elucidate their nature and the infection mechanisms involved. Several years later, the molecular mechanisms underlying these diseases were elucidated and shown to be based on the natural genetic engineering abilities of phytopathogenic soil bacteria of the genus Agrobacterium (Rhizobiaceae family; Chilton et al. 1977, 1980). When various susceptible plants (mainly gymnosperms and dicotyledon species) are wounded in natural ecosystems or fields, they produce simple phenolic substances, such as acetosyringone (Fig. 1), which induces the plasmid-localized vir(virulence) genes of Agrobacterium that are responsible for transferring the T-DNA fragments of the Ti-(tumor inducing) plasmid or Ri-(root inducing) plasmid (in the cases of Agrobacterium tumefaciens and Agrobacterium rhizogenes, respectively) to the plant cells (Nilsson and Olsson 1997). Genes of the transferred DNA fragment mediate the formation of neoplastic crown gall tumors and hairy root tissues, which produce (and secrete into the soil) substances called opines (unusual amino acids; Fig. 1), further used by Agrobacterium as nutrients (Weising and Kahl 1996). Most Agrobacterium strains contain only one T-DNA, but some (like those carrying agropine type Ri-plasmids) transfer two independent T-DNAs, denoted T_L-DNA and T_R-DNA. T_R-DNA has high homology to the T-DNA of the Ti-plasmid

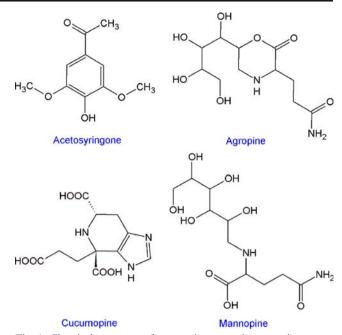
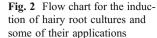
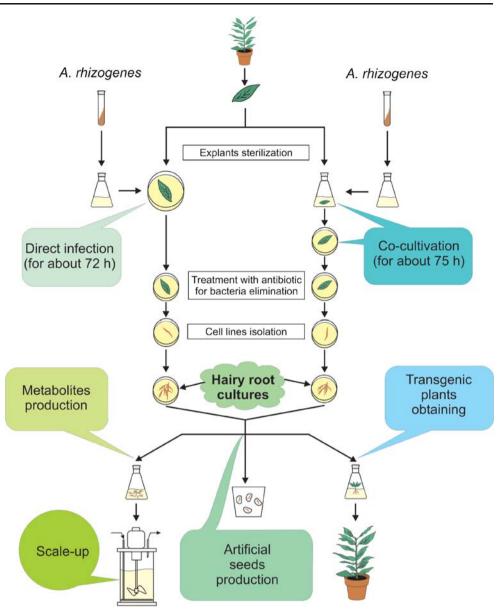


Fig. 1 Chemical structures of acetosyringone and some opines

of A. tumefaciens, while T_I-DNA is strikingly different and has homology to the T-DNA carried by the Ri-plasmid of mannopine A. rhizogenes strains (Nilsson and Olsson 1997). Both T_L-DNA and T_R-DNA are transferred and integrated independently into the host plant genome, but the transfer of T_L-DNA is essential for induction of the hairy root syndrome, and transfer of T_R-DNA does not provoke formation of roots from transformed cultures (Nilsson and Olsson 1997; Sevon and Oksman-Caldentey 2002). T_R-DNA contains two genes, *iaaM* and *iaaH*, responsible for the biosynthesis of auxins and the genes responsible for the synthesis of the opines mannopine (mas1' and mas2') and agropine (ags). T_I-DNA carries 18 open reading frames (ORF), four of which are essential for hairy root formation; ORF10, ORF11, ORF12 and ORF15 corresponding to the genes rolA, rolB, rolC and rolD, respectively. The rolB gene is absolutely essential for the induction of hairy roots. Even when expressed alone, the rolB gene can induce significant hairy root production (Nilsson and Olsson 1997). Detailed information regarding the mechanism involved in the genetic transfer can be found in Gelvin (2000) and Tzfira et al. (2004).

This natural phenomenon is exploited in biotechnology to generate transformed root cultures. The procedure used to induce hairy roots (Fig. 2) includes cultivation of wounded plant parts (called explants) with suspensions of *A. rhizogenes* in aseptic conditions (Sevon and Oksman-Caldentey 2002). Wounded plant explants can be infected with *Agrobacterium* strains either by direct inoculation with bacterial suspensions and incubation on a solid medium or by co-cultivation in liquid media. In either case, the infected explants have to be subsequently transferred to a





solid medium with antibiotics (claforan or penicillin derivates), usually about 72 h later, to eliminate the bacteria (Pavlov et al. 2002; Rahman et al. 2004). The neoplastic, transformed roots (which usually appear after 1-4 weeks) grow in a profusely branched manner, with abundant lateral branches, on Murashige and Skoog or Gamborg's B5 hormone-free media (Pavlov et al. 2002; Sevon and Oksman-Caldentey 2002). Successful genetic transformation can be demonstrated in either of two ways: directly or indirectly, by detecting T-DNA or opines, respectively. The direct way is preferred, as in some cases, opine production is not stable and may even cease (Sevon and Oksman-Caldentey 2002). To detect T-DNA, either polymerase chain reactions (Palazon et al. 2003a; Le Flem-Bonhomme et al. 2004) or Southern blot hybridizations (Nin et al. 1997; Xie et al. 2001) can be used. After a short period of adaptation, the hairy root cultures can then be used to produce metabolites (Wysokinska and Chmiel 1997; Shanks and Morgan 1999; Sevon and Oksman-Caldentey 2002), artificial seeds (Uozumi 2004) or transgenic plants (Choi et al. 2004; Crane et al. 2006). The last two applications are beyond the scope of this review and, thus, will not be further considered.

Many factors influence the likelihood of successful transformation of the mother plant tissue and hairy root induction, including the species, age and type of plant tissue (Sevon and Oksman-Caldentey 2002), the type of *Agrobacterium* strain and the density of the bacterial suspension (Park and Facchini 2000). A number of chemicals may also promote these processes, e.g., acetosyringone (Fig. 1; Joubert et al. 2002). A new technique, named sonication-assisted *Agrobacterium*-mediated transformation (SAAT), to induce the formation of hairy roots in plant species that are difficult to transform, has also been

developed recently. Trick and Finer (1997) observed that the SAAT treatment produces small, uniform fissures and channels in tissues of various plants, which facilitate the access of *A. rhizogenes* to the internal plant tissues. Recently, this technique was also shown to be effective for transforming the hypocotyl of *Papaver* plants by Le Flem-Bonhomme et al. (2004).

Hairy roots—in vitro systems for producing valuable phytochemicals

The high biosynthetic potential of transformed root cultures was largely neglected for years, and the investigations that were performed on them mainly focused on the mechanisms underlying hairy root syndrome. However, during the mid-1980s and early 1990s, there were several investigations on their production of biologically active substances, especially alkaloids (Kamada et al. 1986; Flores et al. 1987), which revealed the biosynthetic capacity of the transformed root systems. Nowadays, hairy root cultures receive more attention as biological matrices for producing valuable metabolites, as they have several attractive features, including high genetic stability (compared to undifferentiated cultures) and relatively fast growth rates (compared to normal roots). Furthermore, growth regulators are not required for their cultivation, which is an important consideration, as some hormones are toxic (2,4-dichlorophenoxyacetic acid, for example), so their presence in many end products is unacceptable. In addition, use of hairy root systems is advantageous for the production of a number of secondary metabolites that are synthesized in plant roots and then accumulated in aerial parts of the plant (the

naphthoquinone derivative lawsone, for example), as such metabolites are accumulated at very low amounts, or not at all, in undifferentiated or shoot cultures (Shanks and Morgan 1999). Table 1 lists some of the phytochemicals produced by hairy root cultures, according to reports published from 2000 to 2006.

One of the aspects of transformed root cultures that has attracted intense commercial interest is their potential for protein production (Shanks and Morgan 1999). The proteins they can produce include medicinally important enzymes such as superoxide dismutase (EC 1.15.1.1; Hyeon and Yoo 2002) and peroxidase (EC 1.11.1.7; Kim and Yoo 1996), other enzymes such as phytase (EC 3.1.3.8; using genetically engineered sesame hairy roots; Jin et al. 2005) and foreign proteins, e.g., monoclonal antibodies (Sharp and Doran 2001) and ribosome-inactivating proteins (Thorup et al. 1994). Recently, Doran (2006) reviewed the current status and problems associated with the production of foreign proteins by hairy root cultures (such as low accumulation levels, instability in the culture medium, etc.) and outlined strategies to minimize their degradation and losses. Although some of these problems have not yet been fully resolved, there is little doubt that these applications will continue to expand in the future.

In addition to proteins and well-known, standard metabolites of the mother plant, hairy roots are also potential sources of new, natural compounds. For example, Li et al. (1998) isolated a new compound named licoagrodione from *Glycyrrhiza glabra* cultures, which was shown to possess strong antimicrobial activities, and Berkov et al. (2003) reported the biosynthesis of a new tropane alkaloid ester in tetraploid hairy roots of *Datura stramonium*. Thus, hairy root cultures offer possibilities for

Table 1 Examples of metabolites produced by transformed root cultures^a

Metabolite	Hairy root culture	Biological activity	Reference
Ajmalicine, ajmaline	Rauvolfia micrantha	Antihypertensive	Sudha et al. 2003
Artemisinin	Artemisia annua	Antimalarial	Weathers et al. 2005
Benzylisoquinoline alkaloids	Papaver somniferum; Eschscholzia californica	Analgesic, antibiotic	Park and Facchini 2000
Betalains	Beta vulgaris	Antioxidant, colorant	Pavlov and Bley 2006
Camptothecin	Ophiorrhiza pumila; Camptotheca acuminata	Antitumor	Saito et al. 2001; Lorence et al. 2004
Iridoid glycosides	Harpagophytum procumbens	Anti-inflammatory; analgesic and antidiabetic	Georgiev et al. 2006a
3,4-Dihydroxy-L-phenylalanine	Stizolobium hassjoo	Therapeutic agent against Parkinson's disease	Sung and Huang 2006
Rutin, hispidulin and syringin	Saussurea involucrata	Anti-inflammatory, antifungal	Fu et al. 2006
Scopolamine and hyoscyamine	Datura innoxia	Anticholinergic	Dechaux and Boitel-Conti 2005
Taxol	Taxus brevifolia	Anticancer	Huang et al. 1997
Thiarubrine A	Ambrosia artemisiifolia	Antifungal, antibacterial, antiviral	Bhagwath and Hjortsø 2000
6-Methoxy-podophyllotoxin	Linum album; Linum persicum	Anticancer	Wink et al. 2005

^a Additional examples can be found in recent reviews by Guillon et al. 2006a,b.

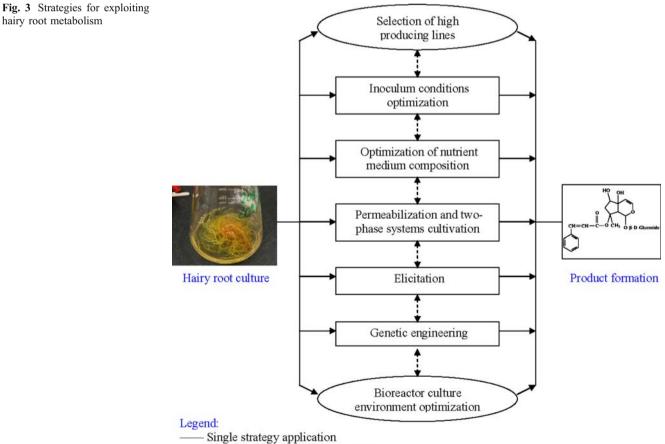
isolating and synthesizing new compounds with potential pharmaceutical value, and several screening programs have been initiated (Kwon et al. 1997).

Strategies for exploiting the secondary metabolism of transformed roots

In order for hairy roots-based technologies to be commercially viable, several criteria have to be met, one of which is to ensure that the yields of the products are acceptable. Because these yields are generally too low at present, several strategies for improving them are being developed, as illustrated in Fig. 3 and discussed below.

Selection Generally, the first step in any attempt to produce a substance from in vitro plant cultures is to select lines that produce as much as possible of it and to continue this selection process after any subsequent optimization steps. Due to the randomness regarding the site at which the T-DNA integrates into the host plant genome, the resulting hairy root cultures often show considerable variations in growth rates, product yields and accumulation patterns of secondary metabolites (Hu and Du 2006). Furthermore, although transformed root cultures are considered to be stable (Shanks and Morgan 1999), a certain amount of heterogeneity has been found in them, and repeated selection seems to be important for obtaining stable, high-productivity hairy root lines (Yukimune et al. 1994). An alternative way for selecting in vitro hairy root lines has been proposed by Flores and Medina-Bolivar (1995), involving the addition of hormones to induce dedifferentiation of hairy roots and subsequent generation of cell suspensions. In the cell suspension stage, well-established approaches for obtaining cell lines that produce high levels of targeted metabolites could be applied (Berlin 1990; Georgiev et al. 2006b), and stable hairy root systems could then be regenerated by simply removing the growth regulators from the medium.

Approaches based on *genetic engineering* represent a relatively new stream of developments. The expression of foreign genes encoding enzymes that catalyze key-limiting steps in the biosynthetic pathways leading to desired metabolites offers good opportunities for improving yields and/or the synthesis of new compounds. Such gene transfers could be comparatively easily performed, using the



----- Combinatorial strategies application

natural genetic engineer (A. rhizogenes), by placing the desired gene(s) between the T-DNA borders and transferring the resulting construct into the host plant genome (Tzfira and Citovsky 2003). As examples of this approach, the gene encoding the enzyme hyoscyamine 6^β-hydroxylase (EC 1.14.11.11), which catalyzes the conversion of hyoscyamine to scopolamine, has been successfully introduced into Hyoscyamus muticus hairy roots (Jouhikainen et al. 1999) and Duboisia hybrid hairy roots. In the latter case, over-expressing plants were also regenerated (Palazon et al. 2003a), and in both cited cases, the genetic manipulations resulted in enhanced scopolamine yields. In another more recent study, Saussurea involucrate was transformed with the chalcone isomerase gene from Saussureamedusa, and the transformed roots accumulated 12 times more apigenin and four times more total flavonoids than the wild-type roots (Li et al. 2006). However, progress in this field is still slow, largely because our knowledge of plant secondary metabolite pathways is limited (Verpoorte et al. 2002). A possible solution to this obstacle is suggested in a recently published work by Rischer et al. (2006), describing attempts to combine genome-wide transcript profiles (using complementary DNA-amplification fragment length polymorphism analyses) with metabolic profiles of elicited Catharanthus roseus cell suspensions. Although substantial further development is required, this kind of gene-tometabolite network analysis seems to be quite promising.

Manipulating the chromosome numbers (*ploidy manipulation*) of in vitro hairy root systems may also provide effective means for both enhancing product yields and the biosynthesis of biologically active substances, which are not found in mother plants. Detailed information about the relationships between ploidy levels and secondary metabolite productivity can be found in Lavania (2005).

Optimization of the inoculum conditions The importance of optimizing the inoculum conditions is frequently underestimated. However, recent investigations have shown that inoculum size and age strongly influence the growth of *Panax ginseng* hairy root cultures (Jeong et al. 2004) and yields of betalains from *Beta vulgaris* hairy root cultures (Pavlov et al. 2003). The first group of cited authors found the growth rate of *P. ginseng* hairy roots to be maximal when a 0.7% (*w/v*) was used and significantly lower when a 0.4% (*w/v*) inoculum was used, and the optimal duration of subculture cycle was found to be 10 days (for *P. ginseng*) and 14 days for hairy roots of *B. vulgaris* (Pavlov et al. 2003; Jeong et al. 2004).

Optimization of nutrient medium composition The nutrient medium also affects hairy roots. Usually, the medium composition is modified with respect to its concentration

(and ratios) of carbon, nitrogen and phosphorous sources (Wilhelmson et al. 2006) and other macronutrients (Sivakumar et al. 2005). A general approach for media optimization is to use statistical analysis (Sung and Huang 2000; Wilhelmson et al. 2006). In the first stage of this approach, the components of the nutrient media (independent variables) are varied, and the resulting culture growth and biosynthesis parameters of the desired metabolite(s) (process outputs) are determined. Multivariate analysis is then applied to explore the interactive "nutrient media-biomass-product" relationships between compounds in the biological systems. Depending on the relationships between the product biosynthesis and root growth parameters, the media can then be either optimized for both growth and production simultaneously or a two-stage (growth followed by bio-production) strategy can be developed.

Elicitation Elicitors are chemicals or biofactors from various sources that can induce physiological and morphological responses and phytoalexin accumulation (Zhao et al. 2005). Hence, elicitors are widely used tools for improving the yields of secondary metabolites that have defense functions in plant cells (Table 2). In recent years, a tendency to exploit abiotic elicitors has developed, as they are generally cheaper than biotic counterparts, and the preparation of some biotic elicitors requires additional facilities for cultivating the source microorganism(s). For example, vanadyl sulfate (VOSO₄), which has been found to be effective in several cases (Bhagwath and Hjortsø 2000; Palazon et al. 2003b), will enhance process costs by about 50 EUR (at concentration 50 mg/l) for production runs in a thousand-liter reactor.

Permeabilization and cultivation in two-phase systems Plant secondary metabolites are generally located in intracellular compartments, usually the vacuoles (Muehlbach 1998). Therefore, the production of many metabolites is limited by the capacity of the vacuole(s). To enhance yields of desired metabolites and reduce total process costs, several approaches for permeabilizing the cells and thereby releasing products have been applied, e.g., treatment with organic solvents (dimethyl sulfoxide; Sim et al. 1994) or surfactants (Boitel-Conti et al. 1996). For instance, treating Datura innoxia hairy roots with Tween 20 led to the migration of significant amounts of alkaloids from cells into the culture medium in a study by Boitel-Conti et al. (1996). Furthermore, the cited authors observed that total alkaloids production (intracellular plus extracellular) was 3- to 8-fold higher in treated than in untreated roots.

The addition of a second, artificial phase can also enhance the yields of target metabolites (Doernenburg and Knorr 1995). Removal of the product(s) by an artificial phase can be especially effective for increasing the total production of products that are subject to feedback

Elicitor	Hairy root culture	Metabolite	Elicitor concentration	Fold increase	Reference
Abiotic elicitors					
CdCl ₂	Brugmansia candida	Hyoscyamine	2 mM	1.35	Pitta-Alvarez et al. 2000
Jasmonic acid	Panax ginseng	Ginsenoside	5.0 mg/l	4.0	Sivakumar et al. 2005
Methyl jasmonate	Panax ginseng	Ginsenoside	22.4 mg/l	4.0	Palazon et al. 2003b
Vanadyl sulfate	Ambrosia artemisiifolia	Thiarubrine A	50 mg/l	8.0	Bhagwath and Hjortsø 2000
Biotic elicitors	-				-
Enterobacter sakazaki	Ammi majus	Umbelliferone	15-ml Suspension/l	1.2	Staniszewska et al. 2003
Fusarium conglutinans	Tagetes patula	Total thiophenes	0.12 mg Carbohydrate/ml medium	2.7	Mukundan and Hjortsø 1990
<i>Phytopthora parasitica</i> (medium filtrate)	Cichorium intybus	Esculin and esculetin	0.1% (v/v)	4.1 and 3.7	Bais et al. 2000
S. aureus	Scopolia parviflora	Scopolamine	133.3 ml Suspension/l	2.8	Jung et al. 2003
Yeast elicitor	Salvia miltiorrhiza	Rosmarinic acid	-	2.3	Chen et al. 2001
Yeast extract	Brugmansia candida	Scopolamine and hyoscyamine	0.8 mg/ml	2.0	Pitta-Alvarez et al. 2000

Table 2 Examples of secondary metabolite yield increases through elicitation^a

^a Additional examples can be found in recent reviews by Guillon et al. 2006a,b

inhibition or degradation by enzymes or other agents. Amberlite XAD resins are widely used as a second phase (Doernenburg and Knorr 1995), partly because many products can adsorb to them non-covalently and non-ionically, facilitating their subsequent elution. Sim et al. (1994) found such use of Amberlite XAD resins (in this case XAD 7) increased both biomass and the accumulation of indole alkaloids in hairy root cultures of *C. roseus*, e.g., catharanthine production was increased to 21.8 mg/l in the presence of the resin from 15.3 mg/l in its absence.

Bioreactors for hairy roots and large-scale cultivation

Bioreactor cultivations represent the final step in the development of techniques for producing metabolites from plant in vitro systems (Bourgaud et al. 2001). Many types of bioreactors have been successfully used for cultivating transformed root cultures (inter alia conventional stirred tanks, stirred tanks with a separate impeller, bubble columns, mist reactors and balloon-type reactors; Curtis 2000). Thus, it would be difficult, if not impossible, to select the "best" bioreactor design for cultivating transformed roots. However, for successful scale-up of hairy root-based processes, whatever type of bioreactor is used, several factors should be considered, including the physiology, morphology, unusual rheological properties and high stress sensitivity of hairy roots (Wysokinska and Chmiel 1997). However, the bioreactor cultivation of hairy roots has been considered in several reviews (Wysokinska and Chmiel 1997; Curtis 2000; Kim et al. 2002). Therefore,

only recent advances (post-2002) will be considered here. One such advance is the development of disposable wave bioreactor systems. The working principle of these systems is based on wave-induced agitation, which significantly reduces stress levels. Moreover, utilization of plastic disposable chambers (called Cellbags) minimizes the need for labor- and time-consuming cleaning and sterilization procedures and facilitates fulfillment of Good Manufacture Practice requirements (Eibl and Eibl 2006). The performance and ginsenoside production of P. ginseng hairy roots in 2-1 wave bioreactors have been studied in detail (Palazon et al. 2003c). The results showed that both biomass accumulation and ginsenoside production were significantly higher in 2-1 wave systems than in shaken flasks. Largescale wave systems with capacities up to 600 l are now commercially available (source: Wave Biotech AG®, Tagelswangen, Switzerland).

Temporary immersion systems (RITA®, Vitpropic, Saint-Mathieu-de-Tréviers, France) have also been used, in our laboratories, for cultivating hairy root cultures from *B. vulgaris* (Pavlov and Bley 2005, 2006) and *Harpagophytum procumbens* (Georgiev unpublished results). Although the RITA® systems have been developed for plant in vitro propagation, their advantages (reduced hyperhydricity and lower consumable and labor costs; McAlister et al. 2005) make them attractive for hairy root cultivation. These systems also seem to be able to solve a problem (so-called hyperhydration), that is associated with the bioreactor cultivation of transformed root cultures in some cases (Georgiev unpublished results). The scale of the RITA® systems is about 200 ml, and the daily cumulative duration of the immersion (flooding) stage can vary from a minute to

several hours. Pavlov and Bley (2006) found that growth of *B. vulgaris* hairy roots is optimal with 15-min immersion/75-min standby cycles, while maximal amounts of betalain pigments are accumulated with 15-min flooding/60-min standby cycles.

Another significant problem that can arise during the cultivation of hairy roots in bioreactor is associated with mass transfer limitations and heterogeneities in root oxygen demands, which are greater in the meristems than the mature tissue (Asplund and Curtis 2001). The hairy roots typically grow in a "tuft-like" manner, which promotes the formation of oxygen and nutrient gradients in the tissue. Curtis (2000) concluded that at low tissue concentrations (<10-g dry weight/l), hairy roots can be grown in bioreactors with virtually any configuration. However, at high tissue concentrations (>10-g dry weight/l) in submerged bioreactors several scale-up limitations may arise. For high tissue density cultivation, Ramakrishnan and Curtis (2004) developed a 14-1 pilot-scale reactor that operates in a "trickle-bed" mode. The dry mass of H. muticus cultivated in this bioreactor reached 36 g/l within 25 days, and the calculated growth index (based on the dry weight) and doubling time were 180 and 3.3 days, respectively, the fastest growth parameters reported for hairy roots so far. In addition, an attempt to cultivate transformed root cultures on a large-scale (500 l) has been reported by Wilson (1997), using a procedure involving inoculum preparation in a 10-1 seed vessel for 2 weeks, followed by aseptic transfer of the root inoculum to the main 500-1 reactor, where the roots were immobilized on barbs, which further facilitate harvesting of the root mass. Although the final biomasses yield (about 4-g dry weight/l) could not be considered high, this procedure addresses many potentially problematic issues (mainly technological) concerned with transferring the tissue inoculum from the seed reactor to the main one(s).

Monitoring the roots' growth during the processes in reactor systems is highly important. Unlike in plant cell suspension-based processes, during the cultivation of organ cultures (e.g. hairy roots), it is impossible to obtain homogeneous samples of the tissue, which complicates measurements of growth and other process parameters. Therefore, conductivity measurements have been widely used to obtain indirect growth estimates (Heyon and Yoo 2003; Pavlov and Bley 2006). Changes in conductivity are due to the cellular uptake of ionic nutrients (NO_3^- anions usually). Linear relationships between increases in biomass and reductions in medium conductivity can be expressed with the equation:

$$(X_{\rm f} - X_{\rm i}) = (C_{\rm i} - C_{\rm f})\beta, \qquad (1)$$

where $X_{\rm f}$ and $X_{\rm i}$ are the final and initial biomass (g), respectively; $C_{\rm f}$ and $C_{\rm i}$ are the final and initial medium

conductivity (mS/cm), respectively; and β (g cm/mS) is a proportional coefficient. However, the cells in the cultures pass through different growth phases (in which rates of uptake of the nutrients vary), and thus, Eq. (1) can only be strictly applied for exact measurements during the exponential growth phase. It is worth noting that, in some cases, such linear relationships between reductions in conductivity and biomass growth increase do not exist (Georgiev et al. 2006a), indicating that the relationships should be individually determined for every culture. The osmolarity of the culture medium offers another possibility for indirect measurements of the root growth. Osmolarity measurements have several advantages over conductivity measurements, as osmolarity takes into account the total number of moles of all solutes present in the medium (Suresh et al. 2001).

Finally, Lee et al. (1998) observed correlations between redox potential (*Eh*) and glucose, cell and ornithine concentrations and proposed that this parameter could be applied for online measurements of *Brevibacterium ketoglutamicum* growth parameters and metabolite biosynthesis. Use of such indicators could also be considered for monitoring hairy roots growth. Huang and Chou (2006) found that redox potential changes during the cultivation of hairy root cultures in a mist trickling reactor reflect the assimilation of NH_4^+ , NO_3^- , sucrose and growth phases.

Future prospects

Recent advances in Agrobacterium-mediated transformation techniques allow hairy root systems to be induced from large numbers of plant species, including rare and threatened medicinal plants, which should contribute significantly to future attempts to preserve global biodiversity and alleviate associated ecological problems. The immense potential of transformed root cultures for biosynthesizing valuable metabolites has been recognized. Furthermore, the advantages of hairy roots make them attractive alternatives to classical technologies and even plant cell suspension cultures. Further enhancement of knowledge regarding plant metabolic pathways and the mechanisms of their regulation in the near future should give us powerful tools for exploiting more fully the biosynthetic potential of hairy roots and for designing new ways to synthesize compounds. All of this suggests that hairy roots technologies are on the threshold of a new era, in which industrial scale bioproduction will be achieved. Of course, several problems must be addressed, which without any doubt will be solved through deeper collaboration between specialists from various relevant fields.

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