

Hairy Root Culture Through *Agrobacterium rhizogenes* for Enhancement of Secondary Metabolites Production in Medicinal Plants: A Review

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Abstract: Plants are a tremendous source for the discovery of new products with medicinal importance in drug development. Several distinct chemicals derived from plants are used in various important ways. Secondary metabolites are economically important as drugs, flavor, dye, pesticides, and food additives. Plants produce the diversity of secondary metabolites which not only plays an important role in adaption according to the environment but also represents an important source of active pharmaceuticals. The possibility of altering the production of bioactive plant metabolites through tissue culture technology is one of the emerging fields of biotechnology to investigate and enhance the production of secondary metabolites. This enhancement through field cultivation has many defects such as slow growth and low and variable yield due to the environmental and biotic factors. Therefore, hairy root culture has been developed as a more efficient alternative biotechnological tool for secondary metabolite synthesis, regardless of environmental, seasonal, and climatic variations. *In vitro* hairy roots formed by genetic transformation have been efficiently utilized for the synthesis of higher levels of flavonoids due to their biochemical and genetic stability as well as their fast growth in media without phytohormones. The focus of the present review is a detailed assessment of research on rhizogenesis in different plants using *Agrobacterium rhizogenes* for the last twelve years particularly for the enhancement of secondary metabolites. The study reveals different techniques involved for rhizogenesis in different plants, compatibility trends of the desired gene, and modifications in the techniques during these years.

Key words: Hairy root cultures, *Agrobacterium rhizogenes*, rhizogenesis, secondary metabolites.

The rationale of the study

Plant secondary metabolites are unique sources for pharmaceuticals, food additives, flavors, and industrially important biochemicals. Accumulation of such metabolites often occurs in plants subjected to stresses including various elicitors or signal molecules. Secondary metabolites play a major role in the adaptation of plants to the environment and in overcoming stress conditions. Environmental factors viz. temperature, humidity, light intensity, the supply of water, minerals,

and CO₂ influence the growth of a plant and secondary metabolite production.

The principle advantage of recent technology is that it may provide a continuous and reliable source of plant pharmaceuticals and could be used for the large-scale culture of plant cells from which these metabolites can be extracted. Plant cell and tissue cultures hold great promise for controlled production of useful secondary metabolites on demand. The current yield and productivity cannot fulfill the commercial goal of plant

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cell-based bioprocess for the production of most of the secondary metabolites. To stretch the boundary, recent advances, new directions, and opportunities in plant cell-based processes are being critically examined. Strategies to improve the production of secondary metabolites must be considered. The productivity of the desired metabolites is limited by the lack of particular precursors. Biotransformation using an exogenous supply of biosynthetic precursors, genetic manipulation, and metabolic engineering may improve the accumulation of compounds.

Agrobacterium rhizogenes (recently revised as *Rhizobium rhizogenes*)⁸⁰ is responsible for hairy root disease in a broad range of dicotyledonous plants and some gymnosperms. *A. rhizogenes*, a Gram-negative soil bacterium, infects plants, adventitious roots called 'hairy roots' are induced from the infected site⁶⁷. This event occurs due to the transfer of the particular DNA region called transfer DNA (T-DNA) comprising the loci between the TR and TL regions of the root-inducing (Ri) plasmid of the bacterium into the plant genome. The basic molecular mechanism of T-DNA trimming from the Ri plasmid, transfer to plant cells, and integration into the plant genome is known, although the functions of several genes on the T-DNA have not yet been elucidated. The hairy roots are aseptically cultured *in vitro* without added phytohormones.

Although as in other fields of science, new technology brought drastic changes in techniques and outcomes of plant cell culture. A problem arose during the *in vitro* production of secondary metabolites from plants due to the incompatibility of the plant with a particular technique. Every plant has its definite genetic composition and stability, thus the outcome of applied technique is not always progressive. It needs to be optimized with different conditions and different bacterial strains to get success.

The objective of this study is to get an overview of the application of elicitors and *Agrobacterium rhizogenes* mediated hairy root culture for the production of secondary metabolites in medicinal plants. Various elicitors have been reported to enhance secondary metabolite production in the plant. In Table 1, the work re-

garding the use of elicitors has been summarized. Data regarding the compatibility of a specific strain of *A. rhizogenes* mediated hairy root culture used for various plants has been recorded for the last twelve years, presented in Table 2. A collective approach has been made to provide the researcher with the information's regarding the outcomes of biotechnological techniques, particularly hairy root culture and application of precursor for the specific plant. This will make easier the path for new researchers.

Trends in enhanced production of secondary metabolites in plants

Plant cell and tissue cultures can be established routinely under sterile conditions from explants, such as plant leaves, stems, roots, and meristems for multiplication and extraction of secondary metabolites²⁶. Strain improvement, methods for the selection of high-producing cell lines, and medium optimizations can lead to an enhancement in secondary metabolite production³⁰. The capacity for plant cell, tissue, and organ cultures to produce and accumulate many of the same valuable chemical compounds as the parent plant in nature has been recognized almost since the inception of *in vitro* technology. The strong and growing demand in today's market place for natural, renewable products has refocused attention on *in vitro* plant materials as potential factories for secondary phytochemical products and has opened the way for new research exploring secondary product expression *in vitro*²⁶.

Addition of precursor for boosted secondary metabolite production:

Production of secondary metabolites can be enhanced in hairy root cultures by the addition of several types of precursors. Enhanced flavonoid production in hairy root cultures of *Glycyrrhiza uralensis* Fisch by combining the over-expression of Chalcone isomerase gene with the elicitation treatment PEG8000 (2 %) alone, yeast extract (YE) (0.1 %) alone, or both of them, and then the total flavonoids were extracted and measured. The results showed that over a culture period of 3 weeks, the wild-type hairy roots, the untreated transgenic hairy roots, and the double-treated

Table 1. Hairy root culture of different plants used with elicitors for increased metabolites Production (courtesy ⁶¹)

Plant	Metabolites	Elicitors	References
<i>Ammi majus</i>	Coumarine, furocoumarin	BION, <i>Enterobacter sakazaki</i>	65
<i>Brugmansiacandida</i>	Hyoscyamine alkaloid	JA	63
<i>Datura stramonium</i>	Tropane alkaloid	Methyl jasmonate	81
<i>Echium rauwolfii</i>	Pyrolizidine alkaloids	Methyl jasmonate Quercetin and Salicylic acid	43
<i>Oxalis tuberosa</i>	<i>Phytophthora cinnamoni</i>	Harmaline, harmine	2
<i>Psoralea corylifolia</i> L.	Phytoestrogenic isoflavones	Chitosan	38
<i>Panax ginseng</i>	Ginsenoside	Chitosan, MeJA, vanadyl Sulfate	55
<i>Pharbitis nil</i>	Umbelliferone, scopole tin, skimmin	CuSO ₄ , MeJA	79
<i>Salvia miltiorrhiza</i>	Transhinone	Yeast elicitor, Ag	22
<i>Scopolia parviflora</i>	Scopolamine	<i>Pseudomonas aeruginosa</i> , <i>Bacillus cereus</i> , <i>Staphylococcus aureus</i>	84
<i>Solanum tuberosum</i>	Sesquiterpine (rishitin, lubimin, phytuberin, phytuberol), lipo-oxygenase	<i>Rhizoctonia bataticola</i> , B cyclodextrin, MeJA	33

JA: Jasmonic acid; MeJA: Methyl jasmonate

transgenic hairy roots accumulated 0.842, 1.394, and 2.838 (g/100 g DW) of total flavonoids, respectively ⁸⁶. Enhanced morphinan alkaloid production was observed in hairy root cultures of *Papaver bracteatum* by over-expression of salutaridinol 7-o-acetyltransferase gene via *Agrobacterium rhizogenes* mediated transformation ⁶². Enhancement of flavone levels was studied through over expression of *Chalcone isomerase* in hairy root cultures of *Scutellaria baicalensis* ⁵².

Knowledge of all the pathways may pave the way in scaling up of hairy root cultures in fermenter through various combinations of elicitors (Table 1). Therefore, using elicitors to the hairy roots can produce an increased amount of secondary metabolites.

***Agrobacterium rhizogenes* mediated hairy root culture**

Agrobacterium species mostly are pathogenic, generating tumor and hairy roots in the plant by the integration of T-DNA containing pathogenic genes encoding phytohormone and opine synthe-

sis enzymes ⁸. Different strains of *A. rhizogenes* showed diverse hairy root induction efficiency. The strains of *A. rhizogenes* that have usually been applied in hairy root induction of medicinal plants include A4, 15834, LBA9402, MAFF03-01724, R-1601, R-1000 and TR105. The *A. rhizogenes* strains differed widely in their ability to induce hairy roots, were reported from *Linum flavum* leaf discs, with the LBA9402 strain being the most efficient ⁴⁰. The choice of *A. rhizogenes* strains for hairy root induction is host-dependent. For instance, although the A4 strain was considered highly virulent and was shown to be highly effective in inducing hairy roots of many plant species, it was not effective in inducing hairy roots from *Linum flavum* leaf discs ⁴⁰.

Ri plasmid of *Agrobacterium rhizogenes*

Virulent strains of *Agrobacterium* contain tumor-inducing (Ti) or Ri plasmids. *A. rhizogenes* contain Ri plasmid (Fig. 1) possessing different gene segments ¹². The transferred DNA (T-DNA) is referred to as the T-region when located on the Ti or Ri plasmid. During infection with *A.*

Table 2. Secondary metabolites produced using hairy root cultures

No.	Name of plant	Name of secondary metabolites	Strain of <i>Agrobacterium rhizogene</i>	References
1	<i>Andrographis paniculata</i>	Andrographolide	MTCC 532	41
2	<i>Glycinemax</i> (L.) Merr	-	K599	13
3	<i>Nicotiana tabacum</i> cv	Stilbenes,(t-Pn and t-Pt)	A4 strain	25
4	<i>Lactuca serriola</i>	Flavonoids and phenolics	AR15834	46
5	<i>Perovskia abrotanoides</i>	Tanshinones	ATCC 15834TR105, and R1000	15
6	<i>Stevia rebaudiana</i>	Steviol glycosides	LBA9402 and A4T	48
7	<i>Mucuna pruriens</i> (L) DC.	-	MTCC 532 & 2364	77
8	<i>Hyoscyamus reticulatus</i> L	Hyoscyamine and scopolamine	A7	47
9	<i>Rauwolfia serpentine</i> and <i>Solanum khasianum</i>	Ajmaline and Ajmalicine	A4 strain	64
10	<i>Datura stramonium</i>	Hyoscyamine	A4 strain	4
11	<i>Artemisia dracunculul</i> L	Hairy root culture	A4	14
12	<i>Agastache foeniculum</i>	Rosmarinic acid	A4 strain	49
13	<i>Withania somnifera</i> (L)	Withaferin A and withanolide A,	R1000	68
14	<i>Berberis aristata</i> DC	Berberin	MTCC 532 and 2364	3
15	<i>Isatis tinctoria</i>	Rutin, Neohesperidin, Budd-leoside, Qu-ercetin	LBA9402	21
16	<i>Psammosilene tunicoide</i>	Saponin	ATCC 15834	75
17	<i>Rubia cordifolia</i>	Anthraquinones	wild culture	88
18	<i>Picrorhiza kurroa</i>	Picroliv	LBA 9402	74
19	<i>Datura metel</i>	Atropine	A4 strain	58
20	<i>Platycodon grandiflorum</i>	Chlorogenic acid (CGA)	strain R1000	71
21	<i>Capsicum annuum</i>	Capsinoids, quercetin	ATCC 43056and ATCC 43057	44
22	<i>Coleus forskohlii</i>	Forskolin	MTCC 2364	56
23	<i>Tripterygium wilfordii</i> Hook. f.	Triptolide and Wilforine	ATCC 15834 and A4	87
24	<i>Gentiana scabra</i>	Gentiopicroside	ATCC 15834	24
25	<i>Fagopyrum tataricum</i>	Rutin and Quercetin	Wild strain	82
26	<i>Valeriana officinalis</i> L	Valerenic acid	A13	70
27	<i>Tribulus terrestris</i> L	β -carboline alkaloids	AR15834 and GMI9534	59

table 2. (continued).

No.	Name of plant	Name of secondary metabolites	Strain of <i>Agrobacterium rhizogene</i>	References
28	<i>Arnica Montana</i> L	Hairy Root Production	LBA 9402.	50
29	<i>Solanum xanthocarpum</i>	Solasodine, Carpesterol	MTCC 532	31
30	<i>Solidago nemoralis</i>	Hairy Root Production	R1000	23
31	<i>Phyllanthus amarus</i>	Genomic and metabolic fingerprinting	ATCC 15834	1
32	<i>Salvia miltiorrhiza</i>	Tanshinone	C58C1	45
33	<i>Decalepis arrayal pathra</i>	2-hydroxy-4-methoxy benzaldehyde (MBALD)	A4,MTCC 532,TR105,and LBA 5402	66
34	<i>Salvia miltiorrhiza</i>	Tanshinone	BCRC15010	20
35	<i>Bacopa monnieri</i> ,	Bacopa saponins	LBA 9402	42
36	<i>Scutellaria viscidula</i>	Flavonoids (Baicalein,wogonoside, wogonin)	Wild culture	39
37	<i>Plumbago indica</i>	Plumbagin	ATCC 15834	19
38	<i>Camptotheca acuminata</i>	Camptothecin	C58C1	78
39	<i>Glycyrrhiza nuralensis</i>	Licochalcone A	ATCC 10060	83
40	<i>Salvia involucrata</i>	Apigenin	Hairy root expressing (CaMV) 35SCHI gene	51
41	<i>Scrophularia buergeriana</i>	E-p-methoxy cinnamic acid	R1000	53
42	<i>Linumtauricum</i>	Arltetraline lignans	ATCC 15834	28
43	<i>Hypericum species</i>	Hairy roots	ATCC 15834and A4	32
44	<i>Linum tauricum ssp.</i>	Lignin	TR 105 andATCC 15834	27
45	<i>Maytenus senegalensis</i>	Hairy roots	LBA9402 and A4T	29
46	<i>Arachis Hypogaea</i> L	Resveratrol	R1601	34
47	<i>Salvia sclarea</i>	Salvipisone	LBA 9402	35
48	<i>Centella asiatica</i> (L)	Asiaticoside	R1000harboring pCAMBIA1302	36
49	<i>Vitis amurensis</i>	Resveratrol	<i>rolB</i> gene of <i>Agro rhizogenes</i>	73

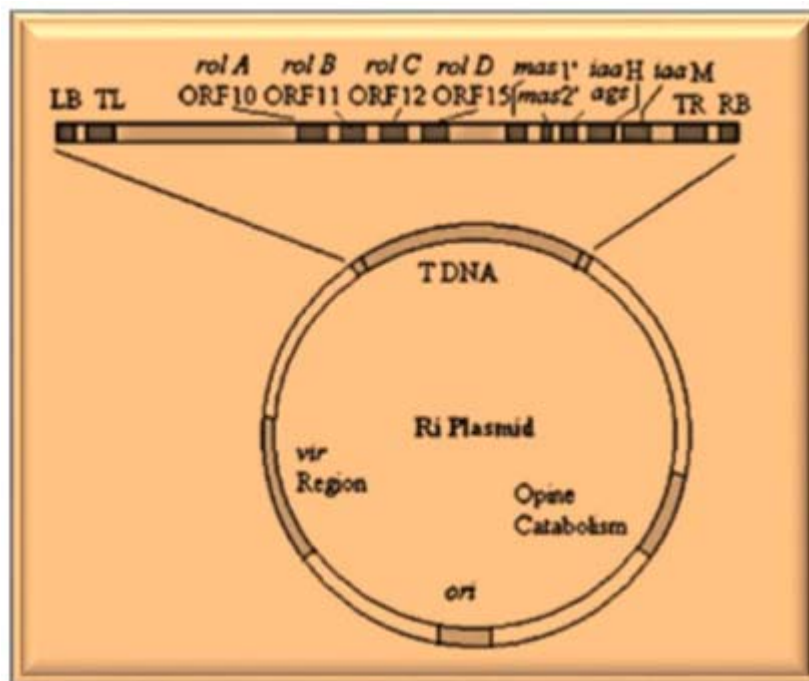


Fig. 1. Schematic representation of Ri plasmid of *A. rhizogenes* (Courtesy ^[12])

rhizogenes, a piece of DNA is transferred from the bacterium to the plant cell. This piece of DNA is a copy of a segment called T-DNA¹⁰. T-DNA is part of the approximately 200 kb Ti/Ri plasmid present in *Agrobacterium* which encodes functions for Ti/Ri plasmid conjugation, opine synthesis, and catabolism and the initiation, transfer, and integration of the T-DNA. T-regions on native Ti and Ri plasmids are approximately 10-30 kbp in size. T-regions are defined by T-DNA border sequences. These borders are 25 bp in length and highly homologous in sequence. They flank the T-region in a directly repeated orientation¹⁷.

Ri plasmids can be classified according to the opines produced. In nopaline Ti plasmid, mannopine, and cucumopine Ri plasmid types, a single T-DNA has been found, whereas, in octopine (pTi) and agropine (pRi) types (Fig. 1), two regions (TL-DNA and TR-DNA) have been identified. Two T-DNAs are separated from each other by about 15 kb of non-transferred DNA. In the central, less conserved region of the TL-DNA of agropine T-DNA, the root-inducing (*rol*) genes are located. TR-DNA contains two genes, *iaaM*, and *iaaH*, responsible for the biosynthesis of auxins¹¹ and the genes responsible for the synthesis of the opines mannopine (*mas10* and *mas20*) and

agropine (*ags*)¹².

Both TL-DNA and TR-DNA are transferred and integrated independently into the host plant genome, but the transfer of TL-DNA is essential for induction of the hairy root syndrome, and transfer of TR-DNA does not provoke the formation of roots from transformed cultures. Detailed information regarding the mechanism involved in the genetic transfer can be referred to in literature^{7,16,76}.

Effect of *rolA*, *rolB* and *rolC* genes in secondary metabolism

Agrobacterium rhizogenes rolA, *rolB* and *rolC* oncogenes have always been considered to be modulators of plant growth and cell differentiation. These *rol* genes are potential activators of secondary metabolism in transformed cells from the Solanaceae, Araliaceae, Rubiaceae, Vitaceae, and Rosaceae families⁶. Studies were made on the activity of *rol* genes individually and their combined action on secondary metabolism⁶⁰.

In transformed plant cell cultures, the *rolC* gene alone can stimulate the production of tropane alkaloids, pyridine alkaloids, indole alkaloids, ginsenosides. The *rolB* and *rolC* gene activate the production of anthraquinones and stilbenes. The

stimulatory effect of the *rolA* gene on nicotine production was also observed⁵⁴. However, *rolA* and *rolB* failed to stimulate ginsenoside production in transformed ginseng calli, and, similarly, the production of caffeic acid metabolites was reduced in *rolC* transformed callus cultures of *Eritrichium sericeum* and *Lithospermum erythrorhizon*⁵.

Studies were also made to know the effect of *rol* genes in secondary metabolites production⁷. In this study, two series of *Atropa belladonna* hairy root lines were obtained: the first transformed via *A. tumefaciens* harboring *rolC* and npt II genes, and the other transformed with *rolABC* and npt II genes. Hyoscyamine and Scopolamine production was measured after 3 and 4 weeks of culture to evaluate the possible role of *rolC* gene in tropane alkaloid formation. The *rolC* gene alone played a significant role (17-fold increase) in the hairy root growth rate. However the *rolABC* genes together led to a much higher (75-fold increase) increase in hairy root growth rate. In contrast, the *rolC* gene alone was as efficient as the *rolABC* genes together (mean value of total alkaloids: 0.36 % dry weight, i.e., 12-fold more than in untransformed roots) to stimulate the biosynthesis of tropane alkaloids in *A. belladonna* hairy root cultures. A correlation exists between the expression of the *rolC* gene and tropane alkaloids, *Catharanthus roseus* alkaloids, and ginsenoside production⁷. Secondary metabolites can be classified based on chemical structure, composition, solubility in various solvents, or the pathway by which they are synthesized. Based on their biosynthetic origins, plant secondary metabolites can be divided into three major groups:

Flavonoids and allied phenolic and polyphenolic compounds

Phenolic compounds confer unique taste, flavor, and health-promoting properties found in vegetables and fruits⁶⁹. Like as phenolic acids, flavonoids are secondary metabolites of plants with polyphenolic structure. They are synthesized by the polypropanoid pathway and the start-up component is phenylalanine molecule. The biological effects of these compounds vary. All flavonoids share the basic C6-C3-C6 structural skeleton, consisting of two aromatic C6 rings (A and

B) and a heterocyclic ring (C) that contains one oxygen atom. Flavonoids are well known for their antioxidant activity. An imbalance between antioxidants and free radicals results in oxidative stress, which will/may lead to cellular damage³⁷.

Terpenoids

These are a large and diverse class of naturally occurring organic chemicals similar to terpenes, derived from five-carbon isoprene units assembled and modified in thousands of ways. Most are multicyclic structures that differ from one another not only in functional groups but also in their basic carbon skeletons. These lipids can be found in all classes of living things, and are the largest group of natural products. About 60 % of known natural products are terpenoids. Terpenoids, such as limonene, myrcene, α -pinene, linalool, β -caryophyllene, caryophyllene oxide, nerolidol, and phytol are flavor and fragrance components common in human diets that have been designated as generally recognized as safe (GRAS) by the US Food and Drug Administration and other regulatory agencies. They display unique therapeutic effects that may contribute meaningfully to the entourage effects of cannabis-based medicinal extracts. Phytocannabinoid-terpenoid interactions could produce synergy concerning the treatment of pain, inflammation, depression, anxiety, addiction, epilepsy, cancer, and fungal and bacterial infections⁵⁷.

Nitrogen-containing alkaloids and sulfur-containing compounds

Alkaloids are among the largest groups of secondary metabolites, being extremely diverse in terms of structure and biosynthetic pathways, including more than 20,000 different molecules distributed throughout approximately 20 % of known vascular plants.

Alkaloids are important chemical compounds that serve as a rich reservoir for drug discovery. Several alkaloids like camptothecin⁸⁹ and vinblastine isolated from natural herbs exhibit antiproliferation and antimetastasis effects on various types of cancers both *in vitro* and *in vivo*.

Hairy root culture

Hairy root culture or transgenic hairy root cul-

tures have revolutionized the role of plant tissue culture in secondary metabolite production. They are unique in their genetic and biosynthetic stability, faster in growth, and more easily maintained. *Agrobacterium rhizogenes*-mediated transformation has several features desirable for the production of secondary metabolites. The rapid and efficient induction of hairy roots from explant tissues in a wide variety of plant species, including medicinal plants, has been reported⁸⁵. These hairy roots are characterized by a high growth rate and high root branching without added phytohormones. Furthermore, they often produce secondary metabolites for a long period, unlike intact roots. For these reasons, switching from undifferentiated cell culture to hairy root culture is considered an attractive alternative for the production of many valuable secondary metabolites that originally accumulated in root tissues.

Using this methodology a wide range of chemical compounds including some important flavonoids has been synthesized. Hairy root cultures of many plant species have been widely studied for the production of secondary metabolites which are useful as pharmaceuticals, cosmetics, and food additives¹⁸. Hairy root cultures represent an interesting alternative to dedifferentiated cell cultures for the production of secondary plant products. Because hairy roots originate from a single plant cell infection by *Agrobacterium rhizogenes*, they are usually considered as genetically stable, in contrast with callus lines. Also, in contrast to dedifferentiated cells, the production of secondary metabolites is not repressed during the growth phase of the culture. Therefore, hairy roots usually produce secondary plant compounds without the loss of concentration frequently observed with a callus or cell suspension cultures. Unlike most secondary plant products in cell suspension cultures, it is possible to get a continuous source of secondary metabolites from growing hairy roots⁹.

Significance of hairy root culture

- Harvesting roots for extracting secondary metabolites can destroy whole plants. Therefore, interest in producing secondary metabolites by developing hairy root culture has been raised.

- Hairy root culture potentially grows faster without needing an external supply of auxins. In certain cases, they do not need incubation under the light.
- Due to their high genetic stability all hairy root cultures are stable in metabolite production.
- Yield in hairy root cultures can be altered by optimizing various factors such as carbon source and its concentration, ionic concentration of the medium, pH of the medium, light, temperature, and inoculums.
- Also utilization of techniques like precursor feeding, cell immobilization, elicitation, and biotransformation of hairy root culture can improve secondary metabolite production.

The present review is highlighting medicinal plants transformed successfully through *A. rhizogenes* to get a high level of secondary metabolites during the last 12 years.

Conclusion and future prospects

Metabolic engineering and biotechnological approaches are in use as an alternative production system to overcome the limited availability of biologically active, commercially valuable, and medicinally important plant secondary metabolite compounds. To date, rapid success has been obtained in exploring the molecular mechanisms of T-DNA transfer, interaction with host plant proteins, their role in plant defense signaling, and integration to plant genome for stable gene transfer for successful plant genetic transformation. T-DNA and corresponding expression of *rol* genes alter morphology and plant host secondary metabolism. Plant transformation technology has now reached a platform of commercial reality. T-DNA and *rol* genes affect plant secondary metabolism. In the recent past, hairy root technology has been significantly improved in different fields: the engineering of secondary metabolism, the increased accumulation and excretion of metabolites after elicitation, the production of therapeutically recombinant proteins, the trapping of biomolecules released in the medium and the scaling-up of the culture process. Hairy roots are easy to grow and to transform. The genetic and biochemical stability of these differentiated cultures and their efficient productivity offer substantial

advantages over cell suspensions. The immense potential of the hairy root system for the production of metabolites and phytoremediation has begun to attract private companies. Soon, hairy roots will provide biotechnologists with powerful tools to reach the precious underground resources of the plant kingdom.

The technique is much useful in the conservation of the plants, especially for those, having root as an important organ bearing metabolites of me-

dicinal value. Root harvesting a major cause of plant uprooting makes the plant rare. The important medicinal substances can be produced using the technique of rhizogenesis in the bio fermenter. So that the plant products can be synthesized commercially and will be easily available for human society. The data and techniques presented in the review must be useful for the optimization of techniques for rhizogenesis and enhancement of secondary metabolite in other important plants.

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