



RECENT ADVANCES IN OPTIMIZATION AND MAXIMIZATION OF ARTEMISININ YIELD FOR ARTEMISININ BASED COMBINATION THERAPY

SUGANDHA TIWARI*¹

¹Dayanand Girls (Post Graduate) College, C.S.J.M. University, Kanpur (U.P), India

ABSTRACT

According to WHO artemisinin based combination therapy (ACT) is the only reliable treatment option for multidrug resistant *Plasmodium falciparum* malaria and to control the spread of the disease under its global malaria eradication programmes. Artemisinin is a sesquiterpene lactone endoperoxide isolated from *Artemisia annua* plants belonging to family Asteraceae. Artemisinin and its derivatives are found to be effective against schistosomiasis, breast cancer, human leukaemia, colon cancer and lung carcinomas. Low yield of artemisinin in *A. annua* plants (0.01-0.8%, in some strains 1.5%) is a serious limitation to meet the annual targets of the drug (approximately 115-135 tonnes) for ACT treatments. The present review covers the biosynthesis of artemisinin and various approaches to increase artemisinin content in *A. annua* plants and new researches on heterologous expression systems.

KEYWORDS : *Artemisia annua*, ACT : artemisinin based combination therapy, *Plasmodium falciparum* malaria, heterologous expression



SUGANDHA TIWARI

Dayanand Girls (Post Graduate) College, C.S.J.M. University, Kanpur (U.P), India

INTRODUCTION

Malarial parasite *Plasmodium falciparum* has become resistant to conventional antimalarial drug chloroquine. Resistance is evolving for sulfadoxine- pyrimethamine that replaced chloroquine as antimalarial therapy. Artemisinin and its derivatives naturally formed in *Artemisia annua* plants, have proved to be effective and safe antimalarial agents, exhibiting excellent activities against *Plasmodium falciparum* parasites that are resistant to commonly used antimalarial drugs. Artemisinin based combination therapies (ACTs) involving two or three drugs are recommended by WHO to control multidrug resistant malaria¹.

The following ACTs are currently recommended by WHO for the treatment of uncomplicated falciparum malaria :

Artemether + lumefantrine,
artesunate + amodiaquine,
artesunate + mefloquine,
artesunate + sulfadoxine–pyrimethamine,
dihydroartemisinin + piperazine.

Artemisinin and its semisynthetic derivatives are believed to kill asexual and sexual intra erythrocytic stages of *Plasmodium* by interacting with the heme discarded by the proteolysis of ingested hemoglobin². Artemisinin and its derivatives are also found to be effective against Schistosomiasis caused by parasite *Schistosoma japonica*, *S. mansoni* and *S. haematobia* causing 1.0 -1.5 million disabilities each year³. Oral administration of artemisinin has also been found to prevent and delay the development of 7, 12-dimethylbenz (a) anthracene (DMBA) induced breast cancer in rat⁴. Artemisinin reacts with iron and forms cytotoxic free radical. It is selectively more toxic to cancer than normal cells because cancer cells contain more intracellular free iron. Covalent tagging of artemisinin to transferrin is reported to increase its selectivity and toxicity

towards cancer cells *in vitro*. Artemisinin-transferrin conjugate significantly retarded the growth rate of breast tumors in rat and could be developed into a potent therapeutic agent of cancer in humans⁴. Artemisinin is also reported to be effective against human leukemia, colon cancer and lung carcinomas^{5,6}.

DISTRIBUTION OF ARTEMISININ IN PLANTS

Artemisia annua L. (annual wormwood, qinghao) belongs to the tribe Anthemideae of the Asteroideae, a subfamily of Asteraceae. The plant is native to China and Vietnam, but now cultivated in many parts of the world. It is mentioned in the Pharmacopoeia of the People's Republic of China⁷ and is widely used indigenous medicinal plant, indicated to relieve fever. The ethnopharmacological applications led to the isolation of artemisinin, as the plant's active principle⁸. *A. annua* is an annual determinate shrub and dies after seed set. It is 1.0 - 2.5 m in height, branches are alternately arranged, leaves are aromatic and dissected and contain biseriolate glandular trichomes that sequester artemisinin and essential oil. At the end of vegetative growth, capitula arise in loose panicles containing numerous central bisexual florets and marginal pistillate florets. Both florets and receptacle bear abundant biseriolate trichomes. Recent evidence indicates that artemisinin is compartmentalized and sequestered in glandular trichomes present in the leaves^{9,10} and flowers¹¹. Highest levels of artemisinin are found in flowers and leaves. Content of artemisinin is highly variable and ranges from as low as 0.01% upto 1.5%¹². Maximum artemisinin content occurs at the time of anthesis, when floral glandular trichomes reach physiological maturity. Artemisinin is not detected in roots, seeds and pollens, structures that do not contain trichomes.

A. *annua* plant

Leaf with trichomes

Inflorescence



ARTEMISININ

Artemisinin is a sesquiterpene (C₁₅) lactone endoperoxide and contains a 1,2,4- trioxane ring structure. The peroxide chemical bond is believed to be essential for its antimalarial activity. Complexation of heme with the peroxide bond followed by electron transfer generates an oxy radical that evolves to the ultimate parasitocidal agent². Artemisinin can be quantified by various analytical procedures including thin layer chromatography, gas chromatography, high performance liquid chromatography with ultraviolet or electrochemical detection¹³, radioimmunoassay and enzyme linked immunosorbent assay¹⁴. Artemisinin and its semisynthetic derivatives such as artemether, arteether, artesunate, artelinic acid and dihydroartemisinin are some of the antimalarials semisynthesized from artemisinin. These are being used as monotherapies or as component drugs in ACTs to treat multidrug resistant *P. falciparum* malaria. The total organic synthesis of artemisinin has been achieved^{15,16}, but is very complicated and low yields are obtained. The plant is still the only valid source of artemisinin¹⁷.

ARTEMISININ BIOSYNTHESIS IN PLANTS

Enzymes of Artemisinin biosynthesis are localized in the two outer apical cells of the

glandular secretory trichomes which comprise of ten cells in total¹⁸. All the terpenes share the common precursors of five carbon (C₅) isoprene units. Terpenes are generated by the condensation of two building block isomers, isopentenyl pyrophosphate (IPP) and dimethylallyl diphosphate (DMAPP)¹⁹. Biosynthesis of IPP and DMAPP takes place by two pathways, i.e., mevalonate (MVA) pathway originating from acetyl CoA in the cytoplasm²⁰ and deoxyxylulose phosphate (DXP)/ methylerythritol phosphate (MEP) pathway originating from glyceric acid -3-phosphate and pyruvate²¹ in the plastids. There exist a crosstalk between the MVA pathway operating in the cytosol and DXP/MEP pathway of plastids^{19, 22}. Three phases of artemisinin biosynthesis are^{23,24}:

Isopentenyl pyrophosphate (IPP) is converted to farnesyl diphosphate by the activity of enzyme farnesyl diphosphate synthase. Conversion of farnesyl diphosphate to amorpha 4-11- diene is the first committed step in the biosynthesis of artemisinin .

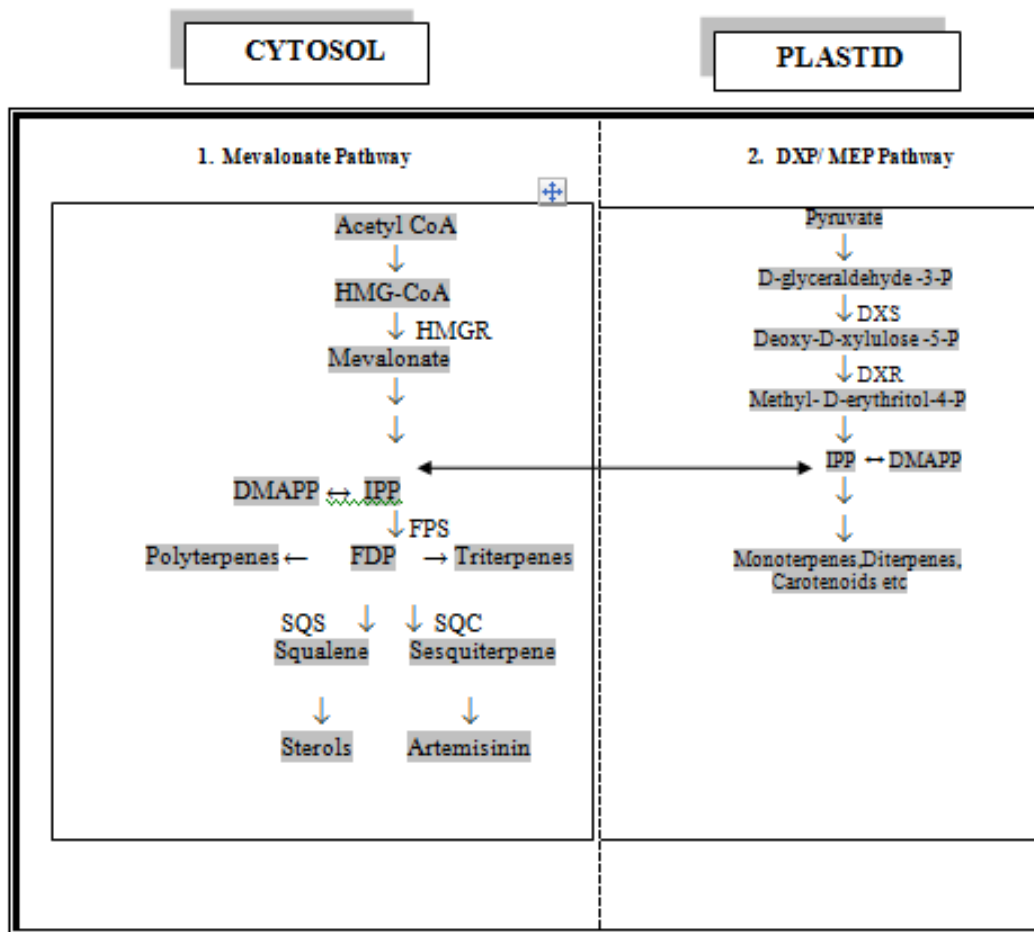
Hydroxylation of amorpha-4,11-diene to artemisinic alcohol is followed by oxidation to artemisinic aldehyde, reduction to dihydroartemisinic aldehyde and then oxidation to dihydroartemisinic acid.

Dihydroartemisinin acid is the true precursor of artemisinin.

process within the plant in the presence of light.

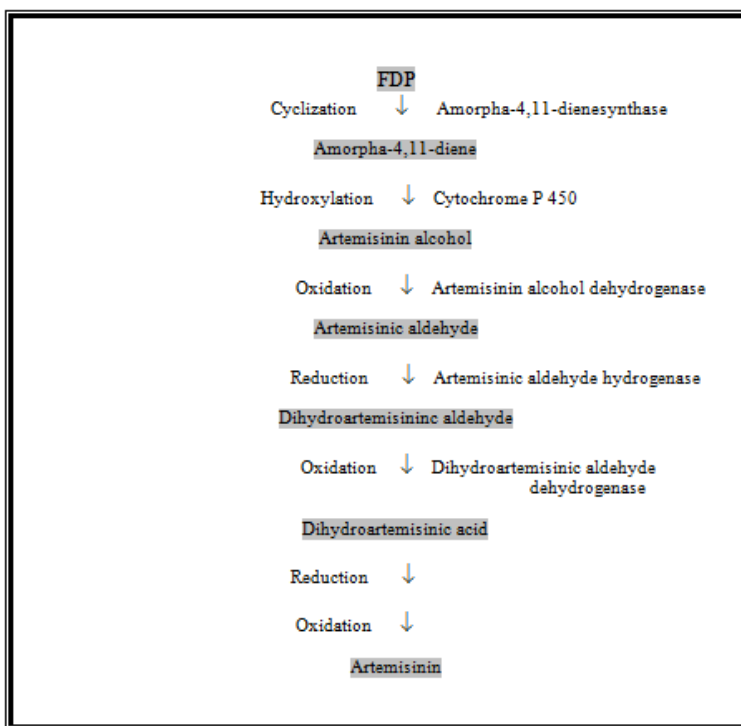
Dihydroartemisinin acid forms artemisinin, probably by non enzymatic **TERPENOID BIOSYNTHESIS PATHWAYS**

(Adapted from Weathers et.al., 2006²⁵)



Some of the regulatory enzymes catalyzing different reactions HMGR, 3-hydroxy-3-methylglutarylcoenzyme A reductase ; FPS, Farnesyl diphosphate synthase; SQC, Sesquiterpene cyclase; SQS, Squalene synthase; DXS, 1-deoxy-D-xylulose-5-

phosphate synthase; DXR, 1-deoxy-D-xylulose-5-phosphate reductoisomerase, IPP, isopentenyl diphosphate; DMAPP, dimethylallyldiphosphate.

PROPOSED PATHWAY FOR THE BIOSYNTHESIS OF ARTEMISININ FROM FARNESYL DIPHOSPHATE (FDP)*(Adapted from Berteau et al., 2005²³)***APPROACHES TO INCREASE ARTEMISININ CONTENT****NON TRANSGENIC APPROACH**

Efforts have been made to increase the content of valuable drug, artemisinin by improving agronomic practices. Several studies have been published on the formation of artemisinin in *A. annua* plants grown in experimental fields. In order to investigate the possibility of isolating greater amounts of the antimalarial compound artemisinin, plants of *A. annua* were cultivated and analysed at different stages of development. Highest content was found just before flowering and it was also possible to correlate development of the plants with the maximum content of artemisinin²⁶. But in Vietnamese plants, highest artemisinin content was found at early developmental stages in five month old plants and not just prior to flowering²⁷. The effect of stage of harvest and plant growth regulators on herbage, artemisinin and essential oil yield of *A. annua* were studied and was found that artemisinin content was

maximum during early flowering stage of growth and declined at later harvests²⁶. Application of GA₃ significantly increased yield and artemisinin content²⁸. Population density of 2.22 plants/ hectare yielded 7.4 Kg of artemisinin and 91 Kg of essential oil in North Central India. The plant architecture and canopy structure presumably favored the synthesis and accumulation of artemisinin resulting in artemisinin rich mature plants²⁹. High artemisinin producing chemotypes appeared to be associated with tall robust plants with long internodes, open branching, dense leaves and thick stems³⁰. Genetic basis of this variation gave evidence of a quantitative inheritance of artemisinin³¹. Additive genetic components were predominant, resulting in a high narrow sense heritability estimate. Thus, good results can be expected from mass selection for the breeding of lines of *A. annua* rich in artemisinin³¹. Dominance variance is also present in the total genetic variability,

indicating that crosses between selected genotypes should generate progenies with particularly high artemisinin content³¹. *Artemisia annua* was introduced in India by Central Institute of Medicinal and Aromatic Plants (CIMAP), Lucknow and four artemisinin rich cultivars have been developed, namely Jeevanraksha, Jeevanraksha 2, Arogya and Jwarharti, the respective artemisinin content in the young leaves is 0.8, 1.0-1.2 and 1.2%. These cultivars are the polycross products of serial selections^{32,33}.

Sequencing of transcriptome of *A. annua* was done to identify genes and markers for fast track breeding. Genetic map was prepared with five linkage groups. Replicated field trials resulted in a quantitative trait loci (QTL) map, that accounted for a significant amount of variation in key traits controlling artemisinin yield. Enrichment for positive QTLs in parents of new high yielding hybrids, confirmed that *A. annua* can be developed into an economically promising crop in terms of better artemisinin yield³⁴. Selection coupled with *in vitro* cloning should be a promising method for crop and product improvement. Relationship of artemisinin content of tissue cultured, greenhouse grown and field grown plants of *A. annua* was analysed and it was found that artemisinin content of the plants grown under long days in the field was highly correlated with the same clones grown under long days in the greenhouse³⁵. The high correlation between artemisinin content for greenhouse and field plants indicates that greenhouse evaluation can provide reliable procedure to select high artemisinin containing plants. In the study artemisinin content only weakly correlated with regenerated clones grown in the greenhouse, or even with tissue cultured clones grown two years apart. This suggests that the weak correlation for artemisinin content between tissue culture and greenhouse plants was a tissue culture effect, perhaps due to somaclonal variation. Therefore, selection for artemisinin from tissue culture clone is not advisable³⁵. Several tissue culture studies were carried out on *A. annua*^{35,36,37} and artemisinin content was found to be highly

variable under tissue culture condition, from trace amounts to 0.063%. Although roots do not produce artemisinin, the presence of roots enhances artemisinin production in shoot cultures³⁸. However, the role of roots in increasing shoot artemisinin is unclear. The possibility that roots may provide a precursor or promoter of artemisinin production that is translocated to shoots needs investigation.

TRANSGENIC APPROACH

Transformed root and shoot cultures of *A. annua* were established by infection with *Agrobacterium rhizogenes* and *A. tumefaciens*. Transformed hairy root cultures of *A. annua* were obtained by co-culture method using leaf segments of *A. annua* and *A. rhizogenes* strain NCIB 8196 or MAFF 03-01724³⁹. The hairy root clones thus obtained grew vigorously on hormone free medium, showing typical transformed morphology. The genetic transformation of the root was proved by opine assay. Artemisinin concentration in *A. tumefaciens* mediated transformed plant was increased to 3.14 fold (31.4mg/g dry wt) by suppressing the expression of SQS (squalene synthase), a key enzyme of sterol pathway that is competitive to artemisinin biosynthetic pathway. This was done by means of RNA mediated RNAi (RNA interference) technique⁴⁰. The study demonstrates that genetic engineering strategy of RNAi could be an effective method of increasing artemisinin content in the plant. Farnesyl diphosphate synthase (FPS) gene and many more genes of artemisinin biosynthetic pathway such as 3-hydroxy-3-methylglutaryl coenzyme A reductase (HMGR), 1-deoxy-D-xylulose-5-phosphate synthase (DXS), 1-deoxy-d-xylulose-5-phosphate synthase (DXPS) and 1-deoxy-d-xylulose-5-phosphate reductoisomerase (DXPR) have been cloned^{41,42,43,44}. Transformed root cultures expressed these genes showing the presence of plastid localized terpenoid biosynthetic pathway⁴³.

PRODUCTION OF ARTEMISININ USING HETEROLOGOUS SYSTEM

During biosynthesis of artemisinin, sesquiterpene synthases form various sesquiterpenes from farnesyl diphosphate. cDNA clones encoding amorpha-4,11-diene

synthase (ADS), a key enzyme in the artemisinin biosynthesis, and epi-cedrol synthase (ECS), a complex sesquiterpene cyclization synthase, were cloned into Cowpea mosaic virus-based viral vector (pEAQ-HT) with Kozak consensus motif and C-terminal histidine tag. The plasmids were transformed into *Agrobacterium* LBA4404 and, agroinfiltrated into *Nicotiana benthamiana* leaves along with vector (pJL3:p19) containing Tomato bushy stunt virus post-transcriptional gene silencing suppressor. The recombinant enzymes ADS and ECS converted farnesyl diphosphate into amorpho-4,11-diene (97 %) and epi-cedrol (96 %) ⁴⁵.

Attempts are in progress to produce artemisinic acid, an intermediate in artemisinin biosynthetic pathway by genetically modifying *Escherichia coli* by transferring the mevalonate pathway genes from *A. annua* to *Escherichia coli*. Synthesis of amorpho-4,11-diene in *E. coli* has been achieved ^{46,47,48}. Heterologous expression of germacrene A synthase in *E. coli* catalysed the cyclation of farnesyl di phosphate to germacrene A ⁴⁹. cDNA clones of epi-cedrol synthase, a sesquiterpene cyclase from *Artemisia annua* L has been isolated. When expressed in *E. coli*, the recombinant epi-cedrol synthase enzyme catalysed the formation of both olefinic (3%) and oxygenated (97%) sesquiterpenes from farnesyl diphosphate ⁵⁰. *Saccharomyces cerevisiae* strains were genetically engineered having mevalonate pathway, amorpho-4,11-diene synthase and a novel cytochrome P 450 monooxygenase (CYP71AV1) from *A. annua*. These engineered yeast strains were capable of producing up to 100 mg/l artemisinic acid. The synthesized artemisinic acid is transported out of the yeast cell and can be purified ⁵¹. Artemisinic acid lacks the endoperoxidebridge, necessary for therapeutic action. Artemisinic acid can be converted chemically to artemisinin in two steps ⁵²: Reduction of the exocyclic double bond ²³ & Photo-sensitized oxidation of the endocyclic double bond to produce the 1,2,4-trioxane ring ²⁴. By overexpressing every enzyme of the mevalonate pathway in

engineered *Saccharomyces cerevisiae* CEN.PK2, production of artemisinic acid was doubled and that of amorpho-4,11-diene was 10- fold higher than artemisinic acid. By standardization of fermentation process for *Saccharomyces cerevisiae* CEN.PK2, > 40g/L amorpho-4,11-diene was produced which was converted to dihydroartemisinic acid and subsequently to artemisinin by a chemical process ⁵³.

Studies are in progress for the optimization of heterologous expression systems and production of artemisinin precursors for which knowledge of complete biosynthetic pathway and its expression in genetically engineered microbe is required.

CONCLUSION

ACT treatment is needed for up to 500 million *P. falciparum* Malaria cases.

There is currently no licenced vaccine against malaria, *in vivo* artemisinin production is low and cost of chemical synthesis is not economically viable. Researches in the field of plant breeding, tissue culture, transgenics and heterologous systems aim to increase artemisinin content in the plants and to increase the supply of artemisinin and its derivatives for cost effective ACT treatments. Metabolic engineering of microbes like *E. coli* and *Saccharomyces cerevisiae* to produce artemisinin is a novel approach for improving the economics of artemisinin production. Synthetic trioxane molecules with endoperoxidebridge are also being tested for their antimalarial potential. Efforts for maximization and optimization of artemisinin yield in heterologous system and synthetic molecules like trioxanes are in infancy and currently all the artemisinin needed for ACT treatment is obtained from field grown *A. annua* plants. Increase in the yield of artemisinin through various approaches would significantly bring down the cost of Malaria treatment.

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