# **Exploring** Agriculture and

# **Biotechnology in tissue culture**

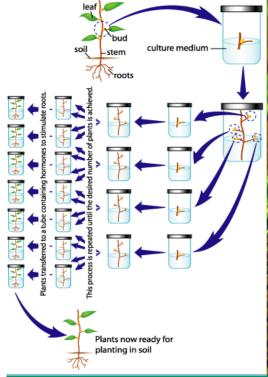


You'll find in-depth information and activities that cover **Tissue culture**, a method of biological research in which fragments of tissue from an animal or plant are transferred to an artificial environment in which they can continue to survive and function. The cultured tissue may consist of a single cell, a population of cells, or a whole or part of an organ. Cells in culture may multiply; change size, form, or function; exhibit specialized activity (muscle cells, for example, may contract); or interact with other cells.



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#### Medical Cannabis and Industrial Cannabis Tissue Culture

The recent legalization of Cannabis in many regions has revealed a need for effective propagation and biotechnologies for the species. Among the planting materials used for cultivation, tissue culture clones provide various advantages such as economies of scale, production of disease-free and true-to-type plants for reducing the risk of GMP-EuGMP level medical cannabis production, as well as the development and application of various technologies for genetic improvement and store germplasm and form the basis for other biotechnologies. Despite this need, research in the area is limited due to the long history of prohibitions and restrictions.

Also, in recent years high-THC (psychoactive) and low-THC (industrial Cannabis) type cannabis has gained immense attention in medical, food, and a plethora of other consumer product markets. Various tissue culture methods have the potential application with cannabis for research, breeding, and novel trait development, as well as commercial mass propagation. Although tissue culture techniques for plant regeneration and micropropagation have been reported for different cannabis genotypes and explant sources, there are significant variations in the response of cultures and the morphogenic pathway. Methods for many high-yielding elite strains are still rudimentary, and protocols are not established. With a recent focus on sequencing and genomics in cannabis, genetic transformation systems are applied to medical cannabis and Cannabis for functional gene annotation via traditional and transient transformation methods to create novel phenotypes by gene expression modulation and to validate gene function. This review presents the status of research focusing on different aspects of tissue culture, including micropropagation, transformation, and the regeneration of medicinal cannabis and industrial Cannabis transformants. Potential future tissue culture research strategies helping elite cannabis breeding and propagation are also presented.

#### Introduction

Cannabis sativa L. is rising to prominence as a commercial crop for industrial, food, medical, and recreational applications. The current wave of interest has been characterized by a growing number of countries easing restrictions around research, commercial cultivation, and sale of dried Cannabis flowers, extracts, and consumable, medicinal, or industrial products. With interest renewed in this crop, which has been cultivated for thousands of years, research and innovation in the coming decades are expected to deepen our understanding of the growth, physiology, and biochemistry of Cannabis. Our improved understanding of this important plant will enable large-scale micropropagation, genetic preservation, and the development of plant biotechnologies for advanced new plant breeding technologies (Bill Carrington, & Marc George Alternative to Plant Characteristics without Traditional Genetic Engineering Molecular/Cell Culture/Tissue Culture https://www.ecronicon.com/ecag/pdf/ECAG-05-00197.pdf ).





As an alternative, *in vitro* techniques offer a promising approach for mass production and germplasm maintenance (Withers and Engelmann, 1997; Watt et al., 2000). Micropropagation can facilitate high throughput propagation in many species and forms the basis of disease-free plants for certified clean plant programs (Lineberger, 1983; Al-Taleb et al., 2011). Tissue culture-based clean plant programs have been used in other vegetatively propagated crops such as potatoes, sweet potato, dates, sugarcane, banana, rice, tobacco, strawberry, grapes, orchids, roses, fruit trees, and some more horticulture of food and ornamental crops, helping to eradicate or prevent the spread of many plant pests, diseases, and viruses (National Clean Plant Network, 2020). Thus, developing an optimized *in vitro* method for propagating clean plants is a crucial strategy to produce large-scale genetically identical plants, retain genetic integrity, and maintain the long-term sustainability of the economically valuable crop (Conway, 2012). This review article aims to provide a comprehensive overview of the most updated available scientific research reported to date on tissue culture in cannabis, to contribute to our understanding of the cannabis tissue culture, and to assess potential applications of the optimized techniques in cannabis plant propagation, regeneration, and transformation.

# Brief History of Cannabis in North America

The relevance of Cannabis as a versatile crop the oilseed, fiber, medicinal, and recreational drug production spans millennia. Between 1000 and 2000 BCE, Cannabis was introduced to Western Asia, Europe, and Egypt as a fiber crop for producing cloth, ship ropes, and paper. After 500 CE, the cultivation of Cannabis was widespread across Europe however, it was not until 1545 and 1606 that it was introduced to South and North America, respectively. Despite its centuries-long cultivation, the beginning of the 20th century saw its recreational use outlawed and medicinal use strongly curtailed by an addendum to the League of Nations' 1912 Opium Convention. This act pushed countries around the globe to restrict and criminalize Cannabis. In Canada, Cannabis was made illegal following its addition to the Opium and Drug Act in 1923 and the United States followed suit with the 1937 Marijuana Tax Act, severely restricting the medicinal use of Cannabis in the United States. Cannabis had been included in the United States Pharmacopoeia since 1850 and was removed in 1942, a few years after passage of the Marijuana Tax Act of 1937. In the United States, Cannabis is classified under the most restrictive drug class (Schedule I) as part of the Comprehensive Drug Abuse Prevention and Control Act of 1970. This 1970 act overturned the 1937 Marijuana Tax Act and states that Cannabis has "no apparent medical potential and a high likelihood of abuse". These restrictions, which made no distinction between fibrous Cannabis and drug-type Cannabis, had the unfortunate consequence of limiting most Cannabis research by making its acquisition for research purposes challenging. Commercial production of industrial Cannabis (Cannabis with 0.3% THC by dry weight) Cannabis in North America, as the distinction between the two, has been largely ignored by government and law enforcement. The strict conditions that regulate Cannabis research have created challenges throughout the research pipeline. Early small-scale.

clinical trials have investigated the use of cannabinoids to treat comorbidities of autism spectrum disorder, anxiety, chronic pain, and seizures and have shown promising results, but research in this area has been highly restricted and progress has been slow. Likewise, these restrictions and the lack of a legal industry have limited research on agronomic, horticultural, and biotechnological aspects of the crop. As a result, relative to the economic importance, technological development is in its infancy and many techniques that are routine for most species are not developed in Cannabis. In recent years, this has started to change as countries around the world have started to lift some restrictions. In Canada, commercial production of Cannabis was legalized in 1998 however, regulatory barriers and a lack of market interest resulted in a very slow-growing industry until recently. In the United States, a pilot-scale production of industrial Cannabis was legalized in 2014 followed by commercial-scale federal legalization in the 2018 farm bill. PrioBefores change, federally funded research in the US could only be conducted with Cannabis obtained from the National Institute on Drug Abuse (NIDA). With the passing of the 2018 farm bill, Cannabis can now be used



for research, but drug-type Cannabis is still highly restricted at the federal level. In 2013, the Marihuana for Medical Purposes Regulations were implemented by the Government of Canada, laying the groundwork for commercial production of medicinal Cannabis. The legalization of the possession, growth, and consumption of Cannabis for recreational purposes followed in October 2018. At the international level, regulations are also beginning to change; a landmark decision by the United Nations Commission on Narcotic Drugs (CND) voted to remove Cannabis from Schedule IV of the 1961 Single Convention on Narcotic Drugs in December 2020, thereby recognizing the medicinal and therapeutic uses of Cannabis. While still highly regulated, the legalization of Cannabis for medical and recreational consumption in Canada, the legalization of Cannabis in the United States, and a similar trend around the world has resulted in a renaissance period for Cannabis research.

#### Industrial Cannabis vs. Medical Cannabis (Marijuana)

Further complicating matters is the legal distinction between Cannabis and drug (narcotic) type cannabis. Any plant containing less than a defined concentration of psychoactive THC is classified as Cannabis. In contrast, anything above the critical limit is classified as drug-type cannabis. Depending upon the jurisdiction, the threshold THC concatenations in flowering plant parts differentiating between industrial Cannabis and drug type cannabis range from 0.2% of dry weight in most European counties, which is 0.3% in Canada, United States, and China and Brazil to 1% in Switzerland, Uruguay, Columbia, Mexico, and several Australian states. While this distinction is not based on taxonomy or genetic relationships, several studies have shown that most Cannabis cultivars are genetically distinct from drug-type cannabis (<u>Rotherham and Harbison, 2011</u>; <u>Cascini et al., 2019</u>). Mainly due to legal restrictions, artificial selection influenced by a decade's long black market, and insufficient knowledge of the *Cannabis* taxonomy, these sub-types are poorly defined, especially the drug type cannabis.

Cannabis is generally cultivated from seed and has named cultivars like most other crops. In contrast, drug type cannabis is generally propagated clonally; the clones are often referred to as 'strains' but are also often referred to as cultivars. As such, any given strain/cultivar can produce various clonal accessions with dramatically different phenotypes, making names unreliable. Further, many strains are offered by different seed companies, and the degree of genetic similarity or difference among providers has not been quantified; therefore, it is generally expected and accepted that there is significant variation within a single strain among seed companies and even within seed lots. Due to these factors, strain names in drug-type cannabis are not reliable regarding a uniform phenotype. *Cannabis indica* and *Cannabis sativa* are the major sources of cannabinoids and are predominantly cultivated, while the third species, *C. ruderalis* is a wild and hardy species and is rarely grown by cultivators as there is no significant content of cannabinoids (Hilling and Mahlberg, 2004). In many lay works of literature, the distinction of 'indica' and 'sativa' have been mentioned and some of the earlier publications have also gathered some phenotypic differences however, there is neither solid taxonomic agreement nor genetic or chemical evidence supporting the differences (Gloss, 2015; Sawler et al., 2015; Piomelli and Russo, 2016). The use of 'indica' and 'sativa' is vaguely based on the general notion that 'Sativa originated from European Cannabis, while 'indica' originated from the Indian subcontinent (Small, 2015), but their exact origin is still debatable.



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#### Table 1: Phenotypic differences among C. indica, Cannabis, and C. ruderalis ecotypes.

| Trait                     | C. indica   | C. sativa   | C. ruderalis  |
|---------------------------|---|---|---|
| Climate                   | Tropical intense sunlight, cool arid<br>regions (Afghanistan, Pakistan,<br>Northern India, Nepal) | Subtropical humid climate, more rainfall,<br>In Mexico, Colombia, Nigeria, Thailand | Northern climates, cool and hilly places, grows in wild (Russia, China) |
| Height                    | Short 1–2 m   | Tall up to 3–4 m  | Very short bushy 0.6–1.0 m  |
| Cannabis female<br>flower | Compact and short inflorescence   | Loose packed and long inflorescence   | Small, compact, very short<br>inflorescence                             |
| Habit                     | Shorter internode   | Longer internode  | Very short internodes   |
| Leaves                    | Broad   | Narrow  | Smaller leaves  |
| Leaf color                | Dark green  | Light green   | Dark leaves   |
| Stalk                     | Shorter woody   | Taller, fibrous   | Short fibrous   |
| Maturity                  | Early maturity 2–3 months   | Late maturity 4–6 months  | Very early maturity 1.5–2 months, autoflowering                         |
| Root system               | Condensed root system   | Deep, expansive   | Shallow smaller   |
| Cannabinoid<br>content    | Lower THC, could be higher CBD  | High THC, Lower CBD in general  | Low THC and CBD   |
| Effect                    | Relaxing effect, inflammation reduction (Medical use preferred)                                   | Incite euphoria, head high (stress relief, recreational use preferred)              | Not grown commercially, only for breeding earliness                     |

Information derived from Schultes et al. (1974); Small and Cronquist (1976), Hillig (2004); Clarke and Merlin (2013), Farag and Kayser (2017), and Small (2017).

# **Figure 1:** Cannabis leaf showing morphological differences of the three different species (*C. indica, Cannabis,* and *C. ruderalis*).



C.indica

C. sativa

C.ruderalis



## **Traditional Cloning in Cannabis**

For decades, seed propagation in cannabis has supported agricultural needs and facilitated genetic improvement. However, with modern horticultural practices in the cannabis industry, stem cutting or traditional cloning, and *in vitro* propagation of this high-value crop has become common practice (Lata et al., 2009a,b, 2011; Potter, 2009). Other methods of propagation are the encapsulation of axillary nodes in calcium alginate beads (Lata et al., 2009a), leaf-derived callus (Lata et al., 2010c), and temporary immersion bioreactor systems (Lata et al., 2010b) but these are limited in lab experiments only. Traditional cloning involves taking stem cuttings from a healthy mother plant and providing a rooting environment for the newly cut clone. For selection as a donor, a clear indication of alternating branches with no visible sign of insects, fungus, or any mineral deficiency in a mother plant is required. Cuttings can be taken from any part of a donor; despite some suggestions that growth in the lower half is better, no difference was observed between cuttings taken from the upper and lower part of the plant (Caplan et al., 2018). However, further research is warranted to test this across more genotypes and conditions. In general, cannabis propagates readily from stem cuttings even without rooting hormones.

Stem cuttings have advantages over seed propagation, including quicker maturation, true-to-type plants, and elite genetics maintenance. Along with the ease of propagation, the practice can limit unwanted gene flow (<u>McKey et al., 2010</u>), for example, between the Cannabis and drug-type, potentially retaining the proportions of active metabolites.

**Figure 2:** Cannabis nodal cloning. (A) Cannabis plants at 6–8 leaf stage. (B) Elongated lateral branches after terminal buds removed from female plants (C) lateral branches planted in the soil after excision from mother plants and. (D) Vegetative clones transferred to 7-inch pots after roots were established and grown. (E) Vegetative clone at maturity.





#### Table 2: Comparison between tissue culture cloning, manual cloning, and seed propagation in cannabis.

| Propagation<br>system        | Seed  | Traditional cloning   | In vitro   |
|------------------------------|---|---|--|
| Roots                        | Tap root is prominent, grow deep, suitable for field cultivation  | Adventitious roots grow from stem laterally, suitable for indoor cultivation  | Adventitious roots grow from stem<br>laterally, suitable for indoor cultivation  |
| Genotype                     | In hybrids, genotype is different for each<br>seed.<br>In feminized seeds, genotype is close to<br>each other               | Same as mother plant  | Same as mother plant   |
| Rooting hormone              | Not required  | 0.1% Indole-3-butyric acid (IBA) is used to<br>promote rooting  | 0.1% IBA is used to promote rooting  |
| Sexual type                  | Segregate in male and female (about 50%<br>each in case of hybrid seeds);<br>near 100% female in case of feminized<br>seeds | All female but chances of developing<br>hermaphrodites or mutated males   | All female but chances of developing<br>hermaphrodites or mutated males  |
| Preferred growing            | Outdoor   | Indoor, hydroponic, aeroponic<br>18:6 h photoperiod   | Indoor/hydroponic/aeroponic<br>18:6 h photoperiod  |
| Preferred Light<br>Condition | Variable between 500 – 2500 $\mu$ molm <sup>2</sup> s <sup>-1</sup>   | Variable between 200 and 300 $\mu molm^2~s^{-1}$  | Variable between 50 and 100 $\mu molm^2$ $s^{-1}$  |
| Yield                        | 500–600 g/plant, relatively long growing<br>cycle and high vegetative growth  | 40–60 g/plant, relatively small plant, short<br>growing cycle, flower matures within<br>2 months  | 40–60 g/plant, relatively, small, short<br>growing cycle, flower matures within<br>2 months  |
| THC%                         | <0.3% THC;<br>mostly used for propagating industrial hemp   | Between 4 and 30% THC depending on strain   | Between 4 and 30% THC depending on strain  |
| Growing medium               | Soil/compost  | Compost/vermiculite<br>cubes/rockwool/hydrotone clay balls  | Sterilized tissue culture medium   |
| Clone health                 | Chances of seedling infection with mites,<br>sucking pests, powdery mildew,<br>Hop latent viroid (Dudding disease)          | Lower chances as grown under controlled<br>condition but could carry disease or pests if<br>cutting come from infected mother plants. If<br>mother plant was infected or symptomless<br>carrier for Hop latent viroid (Dudding<br>disease), chances to carry it forward | Lowest chances as grown under clean<br>condition to carry disease or pests as<br>multiplied from clean stock. Opportuniti<br>to clean for Hop Latent virus as coming<br>from nodal clone stocks free of Hop<br>latent viroid (Dudding disease) |
| Storage                      | 2–3 years in cool dry place   | In a dome for a week  | For a week in controlled condition and<br>up to 12 months at 4°C   |
| Storage<br>requirements      | Protective cover from high sunlight,<br>temperature, and wind; watering as<br>necessary                                     | Cuttings require 65–75% relative humidity;<br>20–23°C temperature and artificial light for<br>growth; proper ventilation  | Controlled and clean purified air HEPA<br>filtered air in culture rooms; 45–50%<br>relative humidity in culture rooms;<br>20–22° C temperature, effective<br>ventilation   |
| Multiplication rate          | One plant can yield thousands of seeds<br>under open pollination/between 100–200<br>seeds from a feminized plant            | 150–200 clones from one month grown<br>vegetative plant   | One to four multiplication rates in one<br>month period but grows exponential in<br>number with time   |
| Hardening<br>requirement     | Not necessary   | About 2–3 days; cuttings are little easier to<br>root and acclimatize in growing environment  | About a week, transition from culture tubes to soil/compost is little riskier  |
| Cost effectiveness           | Can be grown outdoor under little care  | Simpler indoor setup  | Tissue culture lab investment  |
| Preferred use                | Field   | Recreation cannabis   | Medical Cannabis   |

Information derived from Chandra et al. (2008, 2015), Caplan (2018), and Chandra et al. (2020).

On the downside, space for large-scale production is a concern as it can take considerable physical space, representing as much as 20–25% of production space just for cloning. Also, since it is currently manually performed, there is a low multiplication rate, and it is expensive in the long run. Therefore, this technique is more suitable for small growers requiring less than 1000 plants per growth cycle. For this reason, an adaptable, scalable, and robust high throughput tissue culture system with a high multiplication rate that preserves cannabis genetics, and produces more vigorous plants than manual clones, can prove to be more cost-effective in the long run. Even small-scale growers with a small budget use this technique to preserve genetics and test their desired strains' regenerative capacity as a proof-of-concept. Building a team of experts to develop and execute tissue culture protocols successfully can be expensive and time-intensive initially; however, in the long term, it is a promising tool that has benefited many industries, including horticulture and cereal crops (Brown and Thorpe, 1995; Hussain et al., 2012).

Stem cuttings or the traditional cloning method is the widely used propagation system adopted by many growers. *In vitro* propagation is establishing in the cannabis industry slowly and is expected to take over the traditional cloning method. Although stem cuttings and *in vitro* clones can be comparable in terms of vegetative growth and physiological performance (Lata et al., 2009a), *in vitro* clones provide many advantages such as faster multiplication rate, clean clones without disease or virus, cost-effective, etc. Considering these advantages *in vitro* propagation is



expected to become the method of choice for propagation as well as genetic preservation in cannabis soon.

### **Botany and Taxonomy of Cannabis**

Cannabis sativa L. (Cannabis, Cannabis, marijuana) is an annual flowering plant of the family Cannabaceae. Although Cannabis is usually dioecious, hermaphroditism occurs in some cultivars, and both formal and informal breeding programs have resulted in some monoecious cultivars, primarily restricted to Cannabis. The family Cannabaceae consists of ten genera, containing over 100 accepted species, with Humulus lupulus L. (hops; the chief ingredient in beer) being a notable membCannabis isisis native to central Asia, likely in the foothills of the Himalayan Mountain Range. Cannabis is a fast-growing plant, growing up to 10 cm a day and reaching heights of 6 m in its native habitat, while growth in temperate climates is usually lower. When grown from a seed, the first true leaves are pairs of oppositely oriented single leaflets. As the plant matures, the phyllotaxy shifts from opposite to alternate leaf arrange, ment and the number of leaflets per leaf increases (Clarke 1999; Spitzer-Rimon et al. 2019). Leaves on a mature plant are digitate with anywhere from 5 to 11 leaflets and have a long petiole, alth, ough during flowering, they often revert to producing lower numbers of leaflets. Cannabis is predominantly a short-day plant, with flowering induced by 12- to 14-h photoperiods [29]; however, some photoperiod-insensitive cultivars have been developed. Male and female plants cannot easily be distinguished until flowers begin to appear. Male flowers have five green or yellow petals and are larger than female flowers. Female flowers consist of an ovule enclosed in a thin green bract with two yellow/whiteish stigmas emerging from the closed bracts. During the development of the flower, before the elongation of the stigma, glandular trichomes develop on the bract surrounding the ovary.

Two main types of trichomes can be found covering Cannabis plants: glandular and non-glandular trichomes. Only the former produce cannabinoids in any considerable quantity and glandular trichomes are predominantly found on the bracts and floral leaves of female plants. Male plants produce few, if any, glandular trichomes. Due to their low levels of cannabinoids, male plants are generally not consumed as a medicinal or recreational drug and will not be extensively discussed in this review.

**Figure 3:** In Vitro flowering of Cannabis sativa. (A) Flowering Cannabis male plant displaying a hermaphroditic phenotype, showing female flowers (left) adjacent to male flowers (right). Scale bar—1 mm. (B) In Vitro male inflorescences of Cannabis. Scale bar—1 mm (C) A pair of female Cannabis florets obtained from In Vitro flowering Cannabis. Scale bar—1 mm. (D) Glandular trichomes developing on the bract surrounding the ovary of a female Cannabis inflorescence. Scale bar—2 mm. (E) Mature flowering In Vitro explant of Cannabis. Scale bar—1 cm. (F) Four-week-old vegetative explants reverted from In Vitro Cannabis inflorescences. Scale bar—1 cm.

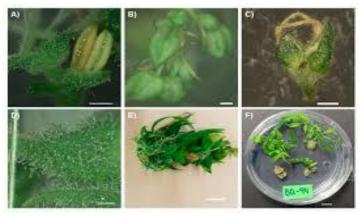




Figure 4: In Vitro germinated seedling of Cannabisdemonstrating opposite leaf arrangement. White arrows show oppositely oriented first true leaves. Scale bar—1 cm. (B) A Stage 2 vegetative explant (subcultured from a nodal explant) of Cannabisdemonstrating alternate leaf arrangement (black arrows), a change in phyllotaxy resulting from explant maturation. Scale bar—1 cm. (C) Cannabisgrown in controlled environment growth chambers under fluorescent lighting. Cannabis grownrown outdoors under shade cloth in Colombia. Image supplied courtesy of Avicanna<sup>™</sup>. (E) Hyperhydric Cannabisexplants growing on Murashige and Skoog (MS) medium supplemented with 0.5 μM TZD TDZ.



The taxonomy of the genus Cannabis is a matter of spirited debate and no consensus has emerged on whether it is a monospecific or polyspecific genus. The ability to distinguish between Cannabis and drug-type Cannabis has been the subject of much interest by law enforcement, which relies on THC content for distinction. Froma law enforcement and regulatory standpoint, the two main categories of Cannabis have been described as "drug-type" (medicinal or recreational) and "fiber-type" (industrial Cannabis), the drug-type generally being dioecious, with a short, wide, bush-like growth pattern, while the fiber-type can be either dioecious or monoecious with a tall and thin growth pattern. However, this distinction is further complicated by Cannabis developed for seed or non-psychoactive cannabinoids, which often morphologically resemble drug-type Cannabis. Two distinct Cannabis chemotypes have been identified, which also fall in line with the two aforementioned morphological groups and are largely defined by their THC content. The fiber-type Cannabis, or "Cannabis", has a THC dry.

weight in the flowering heads of < 1), while the elite drug-type cultivars typically report a THC: CBD ratio >1, or >0.3% THC in the flower heads. However, a taxonomic system based on THC: CBD ratios have faced scrutiny, and other classification systems that further divide the species based on chemotype have been suggested. These include classifications based on other secondary metabolites produced by the Cannabis rather than solely the THC and CBD levels. Early genetic studies attempting to distinguish between the genetic fingerprints of Cannabis and Cannabis have suggested that the chief differentiation factor between the two plants was a single locus that determined the production of THC or CBD synthases. These findings have been echoed by whole genomic and transcriptomic assemblies of Cannabis and drug-type Cannabis, which have shown that Cannabis plants have high levels of



cannabidiolic acid synthase (CBDAS) genes and transcripts, while the THCAS gene encoding the oxide cyclase enzyme, which forms tetrahydrocannabinolic acid (THCA), is dominant in drug-type cultivars. However, recent work using single-nucleotide polymorphisms (SNPs) has shown that the genetic differences between Cannabis extend beyond the loci responsible for cannabinoid production and are instead found throughout the entire genome.

### **Current Utilization and Opportunities for Cannabis Tissue Culture**

Cannabis is a highly adaptable species that can be grown in a variety of conditions, including outdoors in tropical or temperate climates on controlled environments ranging from rudimentary greenhouse structures to sophisticated controlled environment facilities. The production system of choice is determined based on the end-use of the plant. Plants grown to produce low-value commodities such as oilseed or fiber are typically cultivated exclusively outdoors, where production costs are low. In contrast, plants cultivated for dried flowers for recreational or medicinal use can be cultivated outdoors, in greenhouses, on indoor production facilities. While production costs for recreational/medicinal products are also lower outdoors, there is a general belief that indoor production facilities produce higher-quality products, which justifies the extra costs for premium flowers. However, with the growing trend toward extracts and purified cannabinoids likely, at much of the medicinal/recreational production costs. The higher level of oversight offered in controlled environments also allows for easier management of insects and diseases, which is important to meet strict government health and safety regulations surrounding the use of chemical control agents, microbial load, and other quality assurance (QA) requirements. These regulations have driven most of the commercial drug-type Cannabis production into greenhouses and indoor facilities for now.

As with production systems, the approach to plant propagation is influenced by the end-use of the plants. Traditionally, Cannabis has been cultivated by seed using large-scale, highly mechanized, production practices like other grain crops. In contrast, drug/recreational Cannabis is generally propagated using clonal methods and treated as a horticultural crop. This is done to mitigate the high level of phenotypic diversity displayed within seedling populations and to consistently produce high-quality, uniform crops that meet consumer preferences and comply with government regulations. While this variability also exists in Cannabis seeds, the benefits of clonal propagation and manual planting do not justify the costs for oilseed or fiber. However, new regulations surrounding the use of Cannabis to produce CBD and other non-psychoactive cannabinoids have led some Cannabis producers to use clonal propagation.

Clonal propagation can take many forms, but traditionally, Cannabis has been propagated through stem cuttings. In general, Cannabis is relatively easy to root, and large numbers of plants can be produced from a single mother plant. While more expensive than seed, this approach can be efficiently used to mass-produce genetically and phenotypically uniform plants at a commercial scale to produce a more uniform crop. However, this approach requires the maintenance of mother plants in a vegetative state and can occupy 10–15% of the floor space in a commercial operation. The maintenance of mother plants also requires them to remain in a vegetative state. While this is easily accomplished for most genotypes, it presents challenges for day-neutral genotypes as they do not respond to photoperiod. Perhaps of greatest importance, though, is that mother plants are susceptible to insects, pathogens, and viruses and can transmit these biotic factors to their cuttings and lead to problems during production. This is of importance in Cannabis as there are currently very few control options registered for the crop and there is a strong consumer preference for no pesticide use.

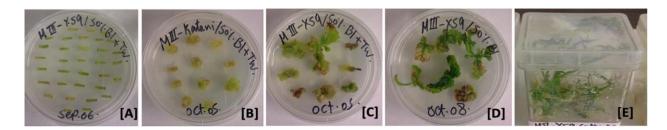


#### In vitro Micropropagation

Although a few cannabis cultivars have regenerated *in vitro Cannabis* spp. have gained a wide reputation for being recalcitrant to tissue culture. At the beginning of the 1970s, along with the conventional propagation system, *in vitro* cultures of cannabis were initiated. The majority of the earlier *in vitro* studies were focused on cannabis callus culture to produce cannabinoids (Veliky and Genest, 1972; Itokawa et al., 1975, 1977; Cannabishill et al., 1978; Hartsel et al., 1983; Loh et al., 1983; Fisse and Andres, 1985). Although there are multiple reports on shoot proliferation via micropropagation there are fewer scientific reports showing regeneration of a full plant through *de novo* regeneration (Mandolino and Ranalli, 1999; Slusarkiewicz-Jarzina et al., 2005; Wielgus et al., 2008; Chaohua et al., 2016).

The majority of regenerated strains and cultivars were monoecious, with few dioecious lines Recently, the optimization of a micropropagation and callogenesis protocol was reported for a few medical cannabis genotypes (Page et al., 2020). Although 48 years passed since thhave e first report of *in vitro* cell culture in cannabis, the available protocols are limited and inconsistent. *In vitro* regeneration of a cannabis plant from a single cell is still a challenge. Thus, the multi-billion-dollar cannabis industry needs an optimized tissue regeneration protocol for both industrial and medical cannabis.

**Figure 5:** Cannabis tissue culture propagation. **(A)** Hypocotyl explants on callus-induction media. **(B)** Hypocotyl explants with the callus-on-callus induction media. **(C, D)** Callus and developing shoots on shoot-induction media. **(E)** Developed shoots on root-induction media.

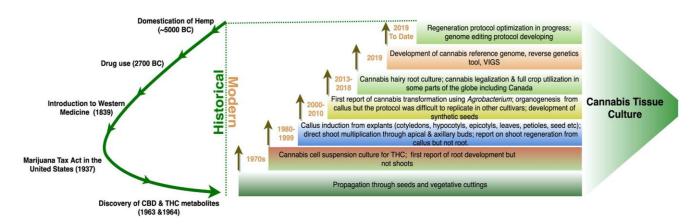


#### Table 3: Cannabis cell culture, transformation, and micropropagation work since 1972–2020.

| Cannable sativa                     | Unknown  | Cell suspension culture for active   | Geeding tissues   | No/No   | Velky and General, 1972                  |
|-------------------------------------|--|--|---|---|--|
| Commentaire receibres               | University   | Assessment of connotancids and<br>assessed of to call a  | Secoling bases  | NewPara   | linkowa et al., 1976                     |
| Carriales Kaliva                    | Chikolowa  | essential of in calca<br>Biotransformation of carriationsia<br>biotransformation carrier   | Choose Riving Lin Lourney   | PalasPala   | Indenious of al., 1877                   |
| Carmable soften                     | Drug type (152 Strein'); Piber type (150   | Carriebhold content In cellus  | Bracts, colys, and leaf   | NewToles  | Hereightli et al., 1070                  |
|                                     | Strain: TUA(2):C-7 1)  |  | tissues   |   |  |
| Connoline radius<br>Cannable sative | Linkonwo<br>Linkonwo   | Root development from callus   | Stondarig   | 2005/240  | Found of Mr. 19811                       |
| Cennets's serve                     | Unknown  | Assessment of metabolites inducing   | Geedling taske  | No/No   | Heltrich and Binder, 1983                |
| Carriable Astros                    | Linkowe  | molitary control to any service or a sublicatory   | Embryo, leat, and stem  |   | Lon et al., 1963                         |
|                                     |  | Diotranaformation of cannabinoid by<br>cell auspenaton culture   | Geedling tasces   | No/No   |  |
| Conversions surfaces                | Livite cours   | Challens michaestern   | More, outplator, and root   | NewTebo   | Fican next Assistant, 11000              |
| Species                             | Gullyara   | Study type   | Explorito type  | Orgonogenesis/Transformation                                      | References                               |
| Carrieble seriles                   | Horney type  | Electing and shoeting from electric<br>outlings  | Astliney absorbs  | WestMan   | Pletter Dumanels et al.,<br>1986         |
| Connolas nativa                     | Linknown and high 1HC/V  | Musiconstantiation of converting   | Londof Tenanalista  | Pdia/Pdia   | Measurement annuk Planin, 1996/          |
| Cannabiz zativa                     | Hemp type  | Preservation procedure of cannabla<br>suspansion cultures  | Florel pert   | No/No   | Jakkel at al., 1989                      |
| Charman dains can alise             | between ge   | Cathor torrestory from of the bod<br>transme   | tread, hypersolyt, solutions,<br>and read   | "We want of the s   | 1000                                     |
| a                                   |  | shoot regeneration from hypocotyl, softwelon, and root   |   |   |  |
| Connebla acens                      | Pedera 10, Pelina 34   | Regeneration of root from calus but no shoot.  | Explant not identified  | Ves/Ves (information not<br>cleacriptive)                         | Madidinnon et al., 2000                  |
| Carnalis satia                      | Interview System   | General transformation and Calue.  | Ethern and leaf   | Parar Visit.  | Fannay and Eurps, 2000                   |
| Cannably patha                      | Citeste (m),<br>Filtrimon-24 (Potential monoecious),<br>Nivrosalisha,<br>Junar 16,<br>Positivar 24 (m) | Full plant regeneration from callus  | Petiole, exiliary bost callus,<br>and callus from internodes  | Ves/No  | **Sluserkiewicz-Jarsha<br>et al., 2005   |
| Closenteedades recedines            | Physiological (1990)<br>Physiological (1990), Physiological statistics (1990)                          | Phorpersonalities of Phoreign  | Photobox, Sourceso, second toboxyour  | "Yea/Max (conty atsolvant is evaluated in the public details are) | Photoconstant of etc. 2008               |
| Consistent nativa                   | Emilotezentee troj, Decisio tou, Stessa troj   | Callus induction and plant reprovation   | Sheep and contribution  | the public detabase)<br>Yea/No                                    | ** Wintputs of al., 2008                 |
| Concernations marthans              | Hornes Lyner   | Hornersonalizers ed abused france enacidatering  | Confidenteurs, informe, soniel reput  | Warms/Pdex  | Commenter and Chronight 201800           |
| Carriabic cativa                    | Hennp type   | Cell suspension culture for secondary<br>metabolism  | Loof Szauna   | No/No   | Flores-Stanches et al., 200              |
| Cermebitr setive                    | MR-1   | Direct organogenesis using nodal<br>segments: synthetic seed development.  | Nodel segments  | Ves/No  | Late et al., 2009a                       |
| Charlestadout matters               | Kimangha   | Street tip suttern   | Statistical Sign  | "Vest-Pace  | Wang of al., SOOR                        |
| Sermately years                     | MRC  | Plegeneration from leaf derived cellus   | Lost torson   | Yes/No  | Late et el., 2010c                       |
| Carriabia sativa                    | N/K  | Synthetic seeds for conservation of<br>clones  | highlat segments  | Ves/Péo   | Lata et al., 2011                        |
| Cermetaly setive                    | Puture77, Delta-llove, Delta405  | Aprobactentum Infection of cannabia<br>roots   | Hypocotyle, cotyledon and<br>cotyledonary node  | Yes/Yes   | Watsby at al., 2013                      |
| Cornalos salva                      | Localization   | Regeneration of plants from callus-  | L-most.   | Yas/No  | ** Humann, 2014 (Bassa)                  |
| Sermebitr vetive                    | Long-me No. 1  | Micropropagation   | Internodea  | Yes/No  | Jiang et al., 2018                       |
| Carenabis Aativa                    | Gridentified   | Caltus industion and Shoot<br>regeneration from collury  | Cotyledon and<br>opicodyledon   | Vinac/Neo   | **Movahadi et al., 2015                  |
| Committe sativa                     | 1. Indeferred fluid  | Cell culture   | Photos  | Pdes/T-Box  | Poreg and Reyser, 2015                   |
| Cavanadoia Antiva                   | Changes  | Full Flant regeneration from callue  | Costylectory  | Yes/No  | Crysonius et al., 2016                   |
| Constantiate seations               | b-Bewerge Types  | Denot requirequirecely to other read and<br>shoot proliferation  | Newford surgrounder   | Monal Teles   | 3 rates set est., 223 540                |
| Caronabie sativa                    | Dialoboteskie and Monica   | Direct organogenesis (shoot and roots)<br>using physicism series.  | Silhoot tipe  | Yes/No  | Grunchova et al., 2017                   |
| Carmabile sativa                    | Webpe  | Direct organogenesis (rooting success<br>of stem cuttings)   | Stem outlings   | Yes/No  | Cepten et al., 2016                      |
| Connectors natives                  | Unknown  | Elementale transformation and<br>regeneration  | Lost expression (for<br>mission and explicit),<br>protoplast (transformation),<br>and polian (transformation) | Unologo/Mag   | Providential of of , PD317<br>(Protects) |
| Controllor, Autors                  | Hemp Landon, Futura, Canda, Jony,<br>CPR 2 and Cherry & Workhamor                                      | Determination of optimal hormons and<br>minurel cells for cellos induction in<br>terms.  | Sillions callingge.   | Winds/Decs  | Thurdon of all, 20138                    |
| Caronable sativa                    | Medicinal cannable but atrain unknown  | Assessment of cannable shoot tips for<br>their models officiency   | Shoot tips and nodal softmas  | Yes/No  | Kodym and Laeb, 2019                     |
| Cermeble setine                     | High THC accessions (1KG2TF, 51525, 155450)  | Regeneration of shoots from immature<br>and mature inforescence  | Ploral thances  | Yes/No  | Plumo at al., 2010                       |
| Dannalais stativa                   | Finals and Explores  | Calue culture, direct regeneration, and<br>pone transformation   | Lowers, policity, and<br>outfory back   | Polar West  | Sectorechicks of al., 2019               |
| Service stolering                   | Tygra, Monoloa, Blaiobrzeakle, Pibrol  | Direct organogenesits (roots) from<br>outlings   | Geedings  | Yea/no  | Simpleatoval et al., 2016                |
| Elementales continen                | 1. Jo that is several  | President and and ad sylence or and almost the feature and another the second s | t cost frames   | Petersters  | Wetton, 2018, 2020<br>(Polent)           |
|                                     | Drug type O. sativa (DA-1)   | Media optimization for callogenesis and  | Leaf and nodal explants   | Yes/No  | Page et al., 2020 (preprin               |
| Cannable sative                     | Hemp, Perimon (m), Felina32 (m), Fedora17  | intercencepalgebiers using september from<br>tastic mode and hereats strates.<br>In vitro plant regeneration and plotdy  |   |   | "Catan-Aulta et al., 2020                |



**Figure 6:** Evolution of cannabis tissue culture research. The green curved arrow on the left shows the key events in cannabis use. Each rectangle on the right shows the major research and development activities at different years. Each brown arrow indicates that the technology is continuously developing, and research work is in progress in the research area.



**Figure 7:** Healthy Cannabis explants growing in We-V boxes (A). Callus cultures growing in glass culture vessels under LED lighting in a controlled environment growth chamber (B) and high-density stackable culture vessels (We-V) with individually programmable LED lighting (C) demonstrate the variety and density with which Cannabis can be cultured under In Vitro conditions.



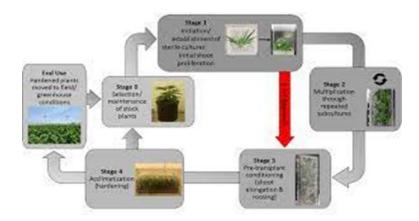
The use of plant tissue culture for the propagation of disease-free plants has provided the foundation for clean plant programs in various crops since the late 1900s. In some cases, certified disease-free plants produced through tissue culture are planted directly in the field for production, while in other cases, they are used as clean material that is further propagated through other means in highly sanitary conditions and tested for important diseases before being used for commercial production. The latter model provides most of the benefits of micropropagation while reducing costs. This approach has been successful in the seed potato industry for developing a disease eradication system. In the case of Cannabis, either approach could be taken, and the decision would need to be based on a careful analysis of the costs and benefits by the producer, which will include many factors such as the efficiency of micropropagation, labor costs, the value of additional floor space, risk assessment, and other factors. The principal challenge in developing effective micropropagation methods is species and genotype specificity, resulting in many variations at each stage of micropropagation. Micropropagation is often broken down into five stages, where each stage needs to



be optimized to establish a fully developed micropropagation method. These include Stage 0: Selection/maintenance of parent plant material; Stage 1: Initiation of cultures; Stage 2: Multiplication of shoots/embryos; Stage 3: Shoot elongation and rooting; Stage 4: Acclimatization. While the selection and maintenance of ex vitro stock plants are often ignored, the importance of stock plant health for the subsequent success of the cultures can have a significant impact on further results. Provided that the stock plants from Stage 0 are in good condition, the explants generally respond well to surface disinfection and produce an initial flush of growth during Stage 1. This initial flush of growth is often followed by a more sporadic growth pattern until the explants acclimatize to In Vitro conditions. It is in Stage 2 after plants acclimatize to In Vitro growth, where the largest benefit of micropropagation becomes apparent: the exponential multiplication of plants. Many horticultural crops are maintained for extended periods of time in Stage 2 and continuously sub-cultured for commercial-scale plant production. To illustrate the capability for rapid plant production, an In Vitro protocol using Stage 2 plants with a reasonable multiplication rate of 10 would produce one million plants after only six subcultures. When enough plants have been produced in Stage 2, they are then transferred to Stage 3 to elongate and develop roots, or alternatively, they are transferred directly from their In Vitro environment to an indoor growth facility/greenhouse to acclimatize, thereby combining Stages 3 and 4. Combining these stages is often preferred for commercial applications as it reduces the number of steps In Vitro, thereby saving time and labor costs. The earliest In Vitro studies of Cannabis were conducted in Cannabis and focused on determining its suitability for In Vitro culture and whether tissue culture would affect the agronomic and chemical characteristics of the plant. Richez-Dumanois et al. showed that Cannabis could be micropropagated using nodal cuttings and the inclusion of IBA and BAP promoted the growth of shoots from existing meristematic tissues. They also demonstrated the successful acclimatization of In Vitro grown Cannabis to greenhouse conditions. Importantly, their work showed that the In Vitro grown plants' chemical and physical profiles were like their greenhouse-grown counterparts. This finding has been reasserted by contemporary studies on

medicinal Cannabis that found micropropagation from nodal cuttings had no significant effect on the cannabinoid contents of the mature flowering plant.

**Figure 8:** The five-stage micropropagation process for tissue culture. The red arrow is indicative of the 1-3-4 approach (where Stage 2 is skipped). Stage 2 is commonly skipped in Cannabis micropropagation methods due to the plant's recalcitrance to long-term culture characterized by a slow decline in fitness. Inclusion of Stage 2 allows for repeated subcultures of In Vitro plants (indicated by the circular arrows), therefore facilitating large-scale multiplication or long-term germplasm storage.







**Table 3:** The table summarizes studies that rely on shoot multiplication (SM) to increase explant numbers. SM refers to the proliferation of multiple shoots from an existing meristem, such as axillary or apical nodes and floral meristems. Cannabis type is defined in this table as either psychoactive "drug-type" (Cannabis; tetrahydrocannabinol (THC) > 0.3% in the flowering head) or known industrial Cannabis genotypes "fiber-type" (Cannabis; THC < 0.3% in the flowering head). A breakdown of the cultivars (CVs) used in the study and the number which responded to the treatment are included for each study. N.S. Not specified is assigned to data that were not specified, instances of "data not shown", or when data are omitted in the original research article.

|   | TBA  | NAA  | LA   | A   | Root  | Root   | Sterra  |
|---|--|--|--|---|---|--|---|
| Treatment   | $(mg(L^{-1}))$   | $(mg_2 L^3)$   | Conge  | $(L^{-3}) = m$  | umber-  | Length   | wide  |
| 1   | 0.1  |  |  |   | 0.65  | +  | ++  |
| 2   | 0.1  | 0.05   |  |   | 2.45  | -+-+-  | +-++-   |
| 3   | 0.1  | 0.25   |  |   | 1.45  | +++-+-   | +-++-   |
| 4   | 0.1  |  | 0.0  |   | 0.80  | -++-   | +-++  |
| 5   | $0.1 \\ 0.5$   |  | 0.3  | 25  | 1.50<br>0.85                                      |  | + + +   |
| 7   | 0.5  | 0.05   |  |   | 0.50  | -  |   |
| -   | 0.5  | 0.25   |  |   | 0.85  | -++-   | + + +   |
| 9   | 0.5  |  | 0.4  | 0.5   | 1.05  | -+-+-  | ++  |
| 1.0   | 0.5  |  | 0.0  | 25  | 1.95  |  | +++-  |
| Table 3. The e  | ffect of di  |  |  |   |   | owth of au   | uxiliary buds   |
| Horme   | ne   | Cone   | centrati   | ion (mg·L   | $\frac{1}{2}$ A                                   | Auxiliary I  | bud number  |
| NA  | A  |  | 0.0  | 05  |   | 1.   | 2 b   |
| NA  |  | 0.   |  | 2.4 a   |   |  |   |
|   | . <b>L</b>   |  |  |   |   |  |   |
| NA  | A  | wed by the s   | 0.<br>ame lette  | -   | significant                                       |  | 1 b<br>at <i>p</i> ≤0.05.   |
| NA<br>Auxiliary bud nu  | A<br>mber follo  | wed by the s   | ame lette  | ers are not   | -   | ly different   | at <i>p</i> ≤0.05.  |
| NA<br>Auxiliary bud nu<br>Table 4   | A<br>mber follo  | -  | ame lette  | ers are not   | ormones   | ly different   | at <i>p</i> ≤0.05.  |
| NA<br>Auxiliary bud nu<br>Table 4<br>Ariation source  | A<br>mber follo<br>. The ana   | lysis of vari<br>SS  | ame lette  | ers are not<br>lifferent h  | ormones<br>F0.05                                  | ly different<br>effect on ro   | at <i>p</i> ≤0.05.<br>o <b>ting.</b>  |
| NA<br>Auxiliary bud nu<br>Table 4<br>Variation source<br>Freatment  | A<br>mber follo<br>. The ana<br>DF   | lysis of vari<br>SS<br>69.95   | ame lette<br>ance of o<br>MS                                   | ers are not<br>lifferent h<br>F Value                               | ormones<br>F0.05                                  | ly different<br>effect on ro<br>Level                                      | at <u>p≤</u> 0.05.<br>oting.<br>F0.01 Level                                 |
| NA<br>Auxiliary bud nu<br>Table 4<br>Variation source<br>Freatment<br>Error   | A<br>mber follo<br>. The ana<br>DF<br>9                                      | lysis of vari<br>SS<br>69.95   | ame lette<br>ance of o<br>MS<br>7.77                           | ers are not<br>lifferent h<br>F Value                               | ormones<br>F0.05                                  | ly different<br>effect on ro<br>Level                                      | at <u>p≤</u> 0.05.<br>oting.<br>F0.01 Level                                 |
| NA<br>Auxiliary bud nu<br>Table 4<br>Variation source<br>Treatment<br>Error<br>Collected total<br>Ta                          | A<br>mber follo<br>. The ana<br>DF<br>9<br>190<br>199<br>ble 5. The          | lysis of vari<br>SS<br>69.95<br>618.65<br>688.6<br>e effect of o                       | ame lette<br>ance of o<br>MS<br>7.77<br>3.26<br>Jifferen       | ers are not<br>lifferent h<br>F Value<br>2.39<br>t media o          | ormones<br>F0.05<br>1.                            | ly different<br>effect on ro<br>Level<br>97                                | at <i>p</i> ≤0.05.<br>oting.<br>F0.01 Level<br>2.59<br>y.                   |
| NA<br>Auxiliary bud nu<br>Table 4<br>Variation source<br>Freatment<br>Error<br>Collected total                                | A<br>mber follo<br>. The ana<br>DF<br>9<br>190<br>199<br>ble 5. The          | lysis of vari<br>SS<br>69.95<br>618.65<br>688.6  | ame lette<br>ance of o<br>MS<br>7.77<br>3.26<br>Jifferen       | ers are not<br>different h<br>F Value<br>2.39                       | ormones<br>F0.05<br>1.<br>on rootin               | ly different<br>effect on ro<br>Level<br>97                                | at <i>p</i> ≤0.05.<br>oting.<br>F0.01 Level<br>2.59<br>y.                   |
| NA<br>Auxiliary bud nu<br>Table 4<br>Auxiliary bud nu<br>Table 4<br>Antion source<br>Treatment<br>Collected total<br>Ta       | A<br>mber follo<br>. The ana<br>DF<br>9<br>190<br>199<br>ble 5. The<br>n Roo | lysis of vari<br>SS<br>69.95<br>618.65<br>688.6<br>e effect of o                       | ame lette<br>ance of o<br>MS<br>7.77<br>3.26<br>Jifferen<br>Ro | ers are not<br>lifferent h<br>F Value<br>2.39<br>t media o          | ormones<br>F0.05<br>1.<br>on rootin<br>Root       | ly different<br>effect on ro<br>Level<br>97<br>g efficienc                 | at <i>p</i> ≤0.05.<br>oting.<br>F0.01 Level<br>2.59<br>y.                   |
| NA<br>Auxiliary bud nu<br>Table 4<br>Variation source<br>Treatment<br>Error<br>Collected total<br>Ta<br>Basic medium          | A<br>mber follo<br>. The ana<br>DF<br>9<br>190<br>199<br>ble 5. The<br>Roo   | lysis of vari<br>SS<br>69.95<br>618.65<br>688.6<br>e effect of o<br>t number           | ame lette<br>ance of o<br>MS<br>7.77<br>3.26<br>differen<br>Ro | ers are not<br>lifferent h<br>F Value<br>2.39<br>t media c<br>oot % | ormones<br>F0.05<br>1.<br>on rootin<br>Root       | ly different<br>effect on ro<br>Level<br>97<br>g efficienc<br>length       | at <i>p</i> ≤0.05.<br>oting.<br>F0.01 Level<br>2.59<br>y.<br>Stem wide      |
| NA<br>Auxiliary bud nu<br>Table 4<br>Variation source<br>Freatment<br>Error<br>Collected total<br>Ta<br>Basic mediun<br>1/2MS | A<br>mber follo<br>. The ana<br>DF<br>9<br>190<br>199<br>ble 5. The<br>h Roo | lysis of vari<br>SS<br>69.95<br>618.65<br>688.6<br>e effect of o<br>t number<br>2.05 a | ame lette<br>ance of o<br>MS<br>7.77<br>3.26<br>lifferen<br>Ro | t media o<br>t media o<br>t 5.0                                     | ormones<br>F0.05<br>1.<br>on rootin<br>Root<br>+- | ly different<br>effect on ro<br>Level<br>97<br>g efficienc<br>length<br>++ | at <i>p</i> ≤0.05.<br>oting.<br>F0.01 Level<br>2.59<br>y.<br>Stem wide<br>+ |

Root numbers followed by the same letters are not significantly different at  $p \le 0.05$  by F test. + Short or thin, ++ Longer or thicker, +++ Longest or thickest



| 9  | 0.5 | _ | 0.05 | 1.05 | ++ | ++ |
|----|-----|---|------|------|----|----|
| 10 | 0.5 | _ | 0.25 | 1.95 | ++ | ++ |

+ Short or thin, ++ Longer or thicker, +++ Longest or thickest

| Hormone | Concentration<br>(mg·L <sup>-1</sup> ) | Auxiliary bud<br>number | Stem<br>wide |
|---------|--|-------------------------|--------------|
| BA      | 1.0                                    | 2.43bc                  | + +          |
| BA      | 2.0                                    | 1.65bc                  | + + +        |
| BA      | 5.0                                    | 1.57c                   | ++           |
| KT      | 1.0                                    | 1.83bc                  | ++           |
| KT      | 2.0                                    | 2.00bc                  | ++           |
| KT      | 5.0                                    | 1.74bc                  | +            |
| TDZ     | 0.1                                    | 2.87ab                  | + + +        |
| TDZ     | 0.2                                    | 3.22a                   | + + +        |
| TDZ     | 0.5                                    | 2.35bc                  | + +          |

#### Table 2 Effect of RA KT and TDZ treatments on plantlet formation

| Auxiliary bud number followed by the same letters are not significantly different at $p \le 0.05$ by F test. |
|--|
| Indie T, - Acialite leterition thile of anticiown incladings isdiated notificanadis canas dented notif       |
| different plant parts grown on media containing organic supplements.   |

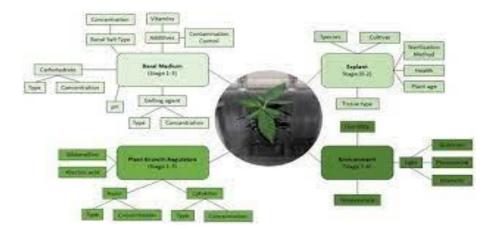
|                    | Organic                  |                      |            |                      |               | Relativ       | ve Retenti           | ion Times                             |                                       |                      |               |                      |
|--------------------|--------------------------|----------------------|------------|----------------------|---------------|---------------|----------------------|---------------------------------------|---------------------------------------|----------------------|---------------|----------------------|
| Callus             | Supplement               | 1                    | 2          | 3                    | CBD<br>(0.74) | CBC<br>(0.76) | 4                    | Δ <sup>8</sup> - <b>THC</b><br>(0.81) | Δ <sup>9</sup> - <b>THC</b><br>(0.84) | 5                    | CBN<br>(0.92) | 6                    |
| Drug 152<br>Bract  | 0<br>1 CH<br>2 CH        | 0.37<br>0.36         | 0.57       | 0.69<br>0.68<br>0.68 |               | _             | 0.80<br>0.79<br>0.80 | _                                     |                                       | 0.91<br>0.91<br>0.91 |               | 0.94<br>0.93<br>0.93 |
|                    | AA 1<br>AA 2<br>NAA-KIN• | 0.37                 | 0.57       | 0.68                 | _             |               | 0.79                 |                                       | _                                     | 0.90                 |               | 0.93                 |
| Calyx              | 0<br>1 CH<br>2 CH        | 0.36                 |            | 0.67                 | -             |               | 0.79                 |                                       |                                       | 0.91<br>0.90<br>0.91 |               | 0.94<br>0.93<br>0.93 |
|                    | AA 1<br>AA 2<br>NAA-KIN• | 0.37<br>0.37         | 0.57       | <br>0.68<br>0.68     |               |               | 0.79<br>0.79<br>0.80 | _                                     |                                       | 0.91<br>0.90<br>0.92 |               | 0.93                 |
| Fiber 150<br>Bract | 0                        | 0.36                 | _          | _                    | _             | _             | 0.80                 | _                                     | _                                     | 0.91                 | _             | 0.93                 |
| Diact              | 1 CH<br>2 CH             | _                    | 0.57       | 0.67                 | -             | _             | 0.30                 |                                       | -                                     | 0.91<br>0.90         | -             | 0.93                 |
|                    | AA 1<br>AA 2<br>NAA-KIN• | 0.38                 | 0.57 ≝<br> | 0.68                 | —             | _             |                      | _                                     | _                                     | 0.91                 | _             |                      |
| Calyx              | 0<br>1 CH                | 0.37<br>0.37<br>0.37 | _          | 0.69                 | -             |               | 0.79<br>0.79         |                                       |                                       | 0.91<br>0.91<br>0.91 |               | 0.93                 |
|                    | 2 CH<br>AA 1             | 0.37                 |            | 0.69<br>0.68         | _             | _             | 0.79<br>0.79         | _                                     | _                                     | 0.90<br>0.91         | _             | 0.93                 |
|                    | AA 2<br>NAA−KIN*         | _                    | 0.57       | 0.68<br>0.69         | _             | _             | 0.79<br>0.80         |                                       | _                                     | 0.91<br>0.92         | _             | 0.94<br>0.94         |

Hormone concentrations: 2,4-D (0.2 mg/l), IAA (1 mg/l), NAA (0.1 mg/l) and kinetin (2 mg/l). \*Medium contained the hormonal combinations (NAA-KIN) which supported optimal growth conditions for each strain as described in table 3. Amino acid (AA) mixtures: No. 1, see table 2: No. 2, Gambor's mixture (23).



As it has been well established that Cannabis can be cultured In Vitro without affecting its biochemical outcomes, contemporary studies have shifted to determining the optimal growth and multiplication conditions for each stage of micropropagation, a task complicated by numerous factors which must be considered when growing a plant In Vitro. Existing Cannabis micropropagation studies have primarily taken to optimizing freshly initiated tissues for shoot proliferation, opting to focus on plant growth regulator (PGR) combinations that result in rapid shoot proliferation. Once developed, the shoots are rooted on an auxin-rich medium and then transferred back into growth facilities (Stages 3 and 4). These rapid and high-throughput approaches to Cannabis micropropagation are useful but neglect to study the long-term health and maintenance of the explants in culture, evidenced by the relatively few studies reporting results from Stage 2. The result of this is that explants are not fully acclimatized to In Vitro conditions, and consequentially, the protocols are only optimized for Stages 1, 3, and 4 (a 1-3-4 approach) rather than for the long-term conservation and multiplication of germplasm in Stage 2.

**Figure 8:** Tissue culture of healthy explants relies on the careful optimization of multiple factors. Center image: Freshly subcultures vegetative explant of Cannabis.



| P    | Startin;<br>plants/<br>ropagatic | vessel |      | plants | plants/ vessel plants/ vess |      |     | Starting with 12<br>plants/ vessel<br>Propagation Factor |      |  |
|------|----------------------------------|--------|------|--------|-----------------------------|------|-----|--|------|--|
| WEEK | 1.5                              | 2      | 4    | 1.5    | 2                           | 4    | 1.5 | 2  | 4    |  |
| 1    | 6                                | 6      | 6    | - 9    | 9                           | 9    | 12  | 12   | 12   |  |
| 5    | 9                                | 12     | 24   | 13     | 18                          | - 36 | 18  | 24   | 48   |  |
| 9    | 13                               | 24     | 96   | 20     | - 36                        | 144  | 27  | 48   | 192  |  |
| 13   | 20                               | 48     | 384  | 30     | 72                          | 576  | 40  | 96   | 768  |  |
| 17   | - 30                             | 96     | 1536 | 45     | 144                         | 2304 | 60  | 192  | 3072 |  |



#### **Genetic Transformation**

An ability to identify, characterize, and apply the genetic variability using biotechnology is the basis of molecular breeding. There are forward and reverse genetics approaches for genetic studies of an uncharacterized allele. With the improvement of sequencing technology, genetic transformation using reverse genetic tools has been an advantage in the molecular breeding program. While cannabis has gained a wide reputation of being recalcitrant to gene transformation and tissue culture, a few reports are describing the methods on gene transformation and regeneration (Feeney and Punja, 2003; Slusarkiewicz-Jarzina et al., 2005; Sirkowski, 2012; Wahby et al., 2013; Schachtsiek et al., 2019). Genome editing holds the potential to develop knockout mutants for significant cannabinoid biosynthesis genes such as THCA synthase, CBDA synthase, and CBGA synthase. Several varieties were tested; most were monoecious, although a few dioecious varieties were also used. In all cases, Agrobacteriummediated gene transfer system was employed and exhibited successful transfer of genes, but the regeneration frequency was low to none. Feeney and Punja (2003) demonstrated the transformation success at the cellular level, but none of their treatments were successful in regeneration. Similarly, Wahby et al. (2013) applied A. rhizogenes strains (A4, AR10, C58, and IVIA251) and could induce hairy roots on the explants derived from hypocotyl and cotyledonary node; however, plantlet regeneration became a bottleneck for them as well. There is two patent information with the claim of successful genome modification and regeneration of cannabis with limited descriptions (Sirkowski, 2012). Thus, there is a need for an optimized protocol for the transformation and regeneration of cannabis replicable and reliable across different species.

#### **Transient Genetic Transformation**

There are various molecular tools developed for transient genetic transformation, including virus-induced gene silencing (VIGS). VIGSis an RNA mediated post-transcriptional gene silencing (PTGS) technique applied to study gene function in a relatively short period (Baulcombe, 1999; Liu et al., 2002; Senthil-Kumar and Mysore, 2014; Adhikary et al., 2019). Once a VIGS protocol is established in a species, it takes 3–6 weeks to see the loss-of-function phenotype of the tested gene/s *in vivo* (Adhikary et al., 2019). Thus, this is an ideal tool to apply, as a proof of concept, to define a target gene's function prior to creating a stable transformation. VIGS, using the Cotton leaf crumple virus (CLCrV), was recently established in *Cannabis*, demonstrating the loss-of-function phenotype of *phytoene desaturase* (*PDS*) and *magnesium chelatase subunit I (ChII)* genes (Schachtsiek et al., 2019). Although the loss-of-function phenotype was weak, the researchers paved a clear path to explore unknown genes' functions in the species. There are viral pathogens reported in cannabis (McPartland, 1996) and many viral vectors developed to date; tobacco rattle virus (TRV) is one of them with a broad-spectrum host range (over 400 plant species) across dicot species (Dinesh-Kumar et al., 2007). Given that TRV can also infect cannabis, potentially demonstrating a strong loss-of-phenotype than CLCrV viral vector.



## **Stable Genetic Transformation**

Both transient and stable transformations have been incredibly beneficial for different research areas and applications in functional genomics. Stable gene transformation is preferred for many applications because once the gene modification is fixed in a plant system, it is heritable. The advantage of the altered gene function can be reaped for generations. As there are numerous reports of successful CRISPR-Cas9 mediated gene editing in many plant species, adopting this newly developed molecular tool in cannabis is vital to improving this economically important plant species. CRISPR can precisely alter a gene's function in a genome (Jinek et al., 2012). It has great potential to benefit both basic and applied plant biology research and development. Therefore, establishing the technology in the cannabis crop is essential for functional studies of thousands of unknown genes and the development of novel varieties.

Traditional genetic modification (GM) and gene editing by CRISPR method are viewed differently (<u>Shew et al., 2018</u>). Gene editing performed using CRISPR method is not considered to be GM organism in some regions. Conventionally, GMO crops refer to organisms that have been altered in a way that they would not have evolved naturally. Moreover, GMO involves transferring foreign DNA fragment from one species to another (transgenic) or within the same species (cisgenic). But in the case of CRISPR edited plants, the targeted mutation is created by using an enzyme and a small guide RNA. While the mutation continues to inherited, the CRISPR machinery can be eliminated in the next generation (<u>Aliaga-Franco et al., 2019</u>). This method is precise and faster than conventional breeding practices, and it is much less controversial than GMO techniques. Therefore, the establishment of CRISPR-Cas9 system in cannabis is another crucial aspect that needs to be explored.

# **Hairy Root Culture**

*Agrobacterium rhizogenes* is another functional genomics tool to assess the function of a gene or developing transgenic plants. These are differentiated cultures that are induced by the infection of *Agrobacterium rhizogenes*, a soil bacterium. Hairy root culture has a high growth rate in a hormone-free medium and exhibits the potential to yield secondary metabolites comparable to the wildtype (Pistelli et al., 2010). It enables the use of stable and reproducible bioreactor-based production and extraction independent of weather conditions, regulatory hurdles, and a lower risk of microbial contamination. This is a safe approach for producing medicinal and active metabolites free of hormones/viruses and does not require pesticides or insecticides. It is also one of the critical avenues for cannabis genetic transformation and functional genomics research.

Calli or hypocotyls infected by *A. rhizogenes* respond with the emergence of hairy roots from the infected site. Hairy roots can be individually selected and tested for a higher production rate of a compound of interest and cryopreserved at –196°C as a pure culture and subculture further for commercial-scale production (<u>Engelmann</u>, 2004). Cannabis hairy root culture has been successfully induced by *A. rhizogenes* (<u>Wahby et al.</u>, 2006, 2013). Hairy root cultures from cannabis callus were also reported using 4 mg/l NAA as a supplement to B5 medium under dark conditions at 25°C (<u>Farag and Kayser</u>, 2015). In the study, the level of THCA and CBDA was less than 2 µg/g dry weight indicating a very low level of cannabinoids present in the hairy root culture under the dark condition with a 28-day growth cycle.



While detectable levels of cannabinoids are not present in *Cannabis* hairy roots, they have been reported to contain choline, atropine, and muscarine (Wahby et al., 2006, 2017). A higher level of these compounds was observed in the *A. rhizogenes* transformed hairy roots compared to non-transformed control. Choline was the most significant compound ranged between 203 and 510 mg/L (control 66–153 mg/L); Atropine with 562–933 µg/L (control 532–553 µg/L); Muscarine with 231–367 µg/L (control undetectable) (Wahby et al., 2017). Additionally, the THCA synthase gene's heterologous expression in tobacco hairy root culture has been successful (Sirikantaramas et al., 2004; Taura et al., 2009).

# **Meristem Culture**

The culture of indeterminate organs, especially the totipotent cells in the apical dome, is a method to obtain many virus clones in a short period (Mori, 1971; Wang and Charles, 1991). The apical dome region has no vascular connection to the developing procambium, leaf primordium, and axillary buds (Wang and Charles, 1991). This lack of vascular connection provides a basis for using the meristem for pathogen elimination as viruses readily travel through the vascular system but do not efficiently transfer from cell to cell. Uninfected cells can be isolated from the meristematic dome (Wang and Charles, 1991; Wu et al., 2020). It is a robust tool for producing virus-free clones that can then be further multiplied at a commercial scale to produce certified virus-free plants. Characteristically, a section of tissue, mostly the apical dome, is dissected either from apical or lateral buds consisting of leaf primordia (no more than 1–2 in number) and apical meristem (0.1–0.5 mm in length) and cultured in a suitable growth medium. Upon induction of the meristem cells under a favorable combination of hormones and growth environment, the cells can continue to develop into a shoot or regenerate into plants through somatic embryogenesis or shoot organogenesis. The regeneration process occasionally gives direct shoot development from the explant, and sometimes morphogenesis occurs indirectly only after the formation of the callus.

There are well-established meristem culture protocols for different model and non-model species (<u>Mori,</u> <u>1971</u>; <u>Mordhorst et al., 2002</u>; <u>Al-Taleb et al., 2011</u>; <u>Spanò et al., 2018</u>), including the closest relative of cannabis, *Humulus lupulus* (Hops), for eliminating virus infection (<u>Grudzinska and Solarska, 2004</u>; <u>Grudzinska et al.,</u> <u>2006</u>; <u>Adams, 2015</u>; <u>Sallie and Jones, 2015</u>). Given the importance of cannabis as a crop, the development of meristem culture for clean plant production could be useful. Unfortunately, this technique is most effective with viral diseases and would not eliminate fungal and bacterial pathogens known to infect cannabis.

# **Protoplast Culture**

For decades, plant protoplasts have been used for genetic transformation, cell fusion, somatic mutation, and more recently, for genome editing (Lei et al., 2015). Significant progress has been made in other crop species in genetic studies using protoplasts; however, for cannabis, studies are in a development phase, with the conditions suitable for the survival of transfected protoplasts and plant regeneration are yet to be optimized. Mesophyll protoplast isolation and transformation of at least three different cannabis cultivars has been reported (Morimoto et al., 2007; Flaishman et al., 2019). Based on the recent study, only about 4% of the protoplasts survived 48 h in liquid culture and plants were not regenerated (Flaishman et al., 2019). Even in the absence of successful regeneration of a whole plant,



protoplasts are of great value in confirming the effectiveness of designed guide RNA (gRNA) prior to their use for the regeneration of gene-edited plants.

#### Somatic Embryogenesis

Somatic embryogenesis is the regeneration of a whole plant from cultured plant cells via embryo formation, from somatic plant cells of various tissues like root, stem, leaf, hypocotyl, cotyledon or petiole (<u>Shen et al., 2018</u>). They morphologically resemble the zygotic embryo's bipolar structure, bear specific embryonic organs, and go through analogous development stages with similar gene expression profiles (<u>Shen et al., 2018</u>). Somatic embryogenesis can occur through direct regeneration. The embryos are developed directly from explant cells, or more commonly through indirect regeneration in which callus develops first, and the development of embryos occurs from callus cells (<u>Sharp et al., 1980</u>).

Plant regeneration via somatic embryogenesis starts with the initiation of embryogenic cultures by culturing various explants on media supplemented with only auxins or a combination of auxins and cytokinins to control cell growth and development (<u>Osborne and McManus, 2005</u>). One exception to this is the use of thidiazuron (TDZ), a cytokinin-like compound that is often used alone to induce somatic embryogenesis (<u>Murthy et al., 1995</u>). The proliferation of embryogenic cultures can occur on solid or in liquid media supplemented with auxins and cytokinins, followed by prematuration of somatic embryos on lower levels of PGRs or PGR free media to stimulate somatic embryo formation and development. Maturation of somatic embryos can occur by culturing on media with reduced osmotic potential or supplemented with abscisic acid (<u>George et al., 2007</u>). This maturation stage is critical for synthetic seed production as it allows embryos to be desiccated, stored, encapsulated, and treated like regular seeds. However, in many somatic embryogenesis systems, the maturation phase has not been developed, and somatic embryos germinate precociously to produce plants.

Somatic embryos are used as a model system in embryology studies; however, somatic embryogenesis's main economic applications are for developing transgenic plants and large-scale virus-free vegetative propagation of elite plant genotypes. The possibility to scale up the propagation using bioreactors has been reported (<u>Hvoslef-Eide and Preil, 2005</u>). Somatic embryos are also ideal for genetic manipulation purposes as they develop from a single cell, thereby reducing the chances of producing chimeric plants, common when relying on shoot organogenesis or shoot proliferation (<u>Dhekney et al., 2016</u>). Other less common uses of somatic embryogenesis include cryopreservation of genetic materials and synthetic seed technology (<u>George et al., 2007</u>).

<u>Feeney and Punja (2003)</u> investigated the somatic embryogenesis and tissue culture propagation of Cannabis. Despite testing various explants and supplements, and variations in the culture medium and changes to the culture environment, there was no successful plantlet regeneration, and a reliable protocol for somatic embryogenesis in cannabis has yet to be published.



# Thin Cell Layer (TCL)

Thin cell layer (TCL) culture utilizes a thin layer of tissue as the explant to allow close contact between wounded cells and nutrients and growth regulators supplied in the medium; this controls the morphogenesis of the cultures (<u>Nhut et</u> <u>al., 2003</u>). This is most useful where larger explants may also contain a high level of endogenous hormones, carbon sources, and other substances that influence and conflict with the effects of exogenous substances placed in the medium and, thus, interfere with development. In general, sterilized TCL explants are excised either longitudinally (0.5–1 mm wide, 5–10 mm long) or transversally (0.1–5 mm thick) prior to culturing (<u>Nhut et al., 2003</u>; <u>Croom et al.,</u> <u>2016</u>). Like other *in vitro* techniques, TCL requires an optimized protocol regarding basal media, PGRs and other added nutrients and growth conditions such as daylength, light intensity, and temperature. These conditions vary for not only the species but can be genotype dependent. It has been widely used in different species, including bamboo, banana, citrus, tomato, rose, *Lilium ledebourii, Bacopa monnieri*, saffron, among others (<u>Nhut et al., 2003</u>; <u>Teixeira da</u> <u>Silva et al., 2007</u>; <u>Mirmasoumi et al., 2013</u>; <u>Croom et al., 2016</u>; <u>Azadi et al., 2017</u>). TCL's potential is yet to be explored in *Cannabis* spp.; however, it may prove to have some utility in the regeneration of genetic transformants in this high value but re calcitrant regeneration crop.

## **Doubled Haploid Production**

Androgenesis is a biological process by which a whole plant regenerates directly from immature pollen (microspores) through the embryogenesis developmental pathway under *in vitro* conditions. While the resulting plant is haploid and inherently sterile, a diploid plant can arise either spontaneously or artificially (<u>Gilles et al., 2017</u>), usually with colchicine, which blocks cytokinesis without blocking chromosome doubling (<u>Galazkajoa and Niemirowicz-Szczytt,</u> 2013). This doubled haploid is homozygous at all loci. Doubled Haploid (DH) plants have been extensively used in plant breeding programs to increase the speed and efficiency with which homozygous lines can be obtained (<u>Alisher et al., 2007</u>). DH technology is traditionally used to genetically stabilize parental lines for F<sub>1</sub> hybrid production. This is important for the rapid integration of new traits through backcross conversion and to develop molecular mapping populations. It is also used to fix desired traits obtained through transformation or mutagenesis and simplify genomic sequencing by eliminating heterozygosity (<u>Ferrie and Mollers, 2011</u>). As such, this technology would be an important tool for both forward and reverse functional genomics studies.

There are two different approaches to develop haploid plants. First, *in situ* methods, using particular pollination techniques such as irradiated pollen, inter-species crosses or so-called 'inducer lines' (<u>Ren et al., 2017</u>); second, *in vitro* methods including the culture of haploid cells (gametes) and their development to haploid embryos and consequently haploid plants through germination. The microspores, which can be harvested in large numbers (millions), are generally isolated for culture as a uniform population. Alternatively, the culture of whole anthers is used to obtain haploid plants through the androgenesis process. The main disadvantage of another culture is the potential for developing a mix of both haploid and diploid plantlets (<u>Elhiti et al., 2010</u>). In this review, we will focus only on the production of doubled haploids from microspores using *in vitro* culture.

One of the most important factors affecting DH production is the microspore developmental stage. It is a complicated factor that has a strong influence on microspore culture's success. It has been reported that only microspores that are



at a stage sufficiently immature have the ability to change their developmental fate from a gametophytic to embryogenic, leading to sporophytic development (<u>Soriano et al., 2013</u>). The most amenable stage is either the uninucleate stage of the microspore or the early binucleate stage, either at or just after the first pollen mitosis. At this developmental stage, the microspore's transcriptional status may still be proliferative and not yet fully differentiated (<u>Malik et al., 2007</u>). Although all microspores within an anther would be roughly of a similar age, not all cells have embryonic competence. Therefore, the incremental differences in the stages of development of individual microspores can be considered significant. To avoid this problem, <u>Bhowmik et al. (2011)</u> introduced a new treatment, discontinuous Percoll gradient centrifugation, to provide a uniform population of *B. napus* isolated microspores at the appropriate stage of development. This approach has consistently produced high embryo yields and consistent embryo development.

#### **Cannabis Microspore Culture**

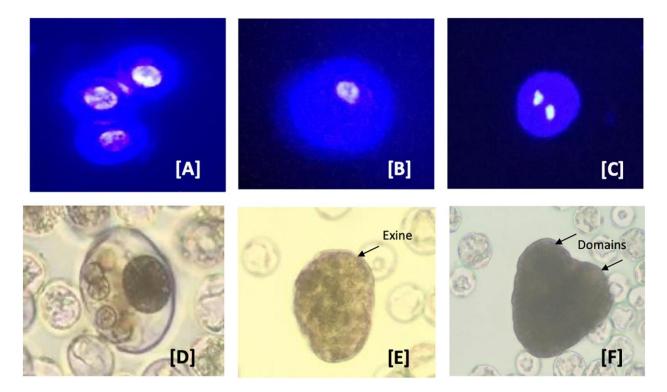
In 2019, an extensive Cannabis breeding program was introduced at Haplotech Inc.<sup>1</sup>. As there has been no previously reported success in the area, a Cannabis DH project was initiated to accelerate this program. Four different Haplotech genotypes were used for this experiment. Both male racemes and pollen-induced female colas were collected, and the buds were fractionated according to size into three groups (2–3, 3–4, and 4–5 mm). Each group was surface sterilized with 15% commercial bleach and washed three times with distilled-sterilized water for 5 min each. The sterilized buds were macerated in isolation media (MS basal fortified by 13% sucrose). The isolated microspores were subjected to fractional centrifugation using Percoll, as described by Bhowmik et al. (2011). The concentration of microspores was diluted to  $4 \times 10^4$  cells/ml with MS basal fortified by 10% sucrose. Five ml of this diluent ( $4 \times 10^4$ ) microspores were mixed with 5 ml of induction media (MS basal, 10% sucrose supplemented with different additives for induction) in 47 mm Petri dishes. The final concentration of the culture used was  $2 \times 10^4$  cells/ml. The isolated microscope.

Samples of isolated microspores were stained with 4, 6-diamidino-2-phenylindole (DAPI) and observed using a fluorescence microscope to monitor their *in vitro* development, once every 3 days. Monitoring of the culture samples by DAPI staining in the first 2 weeks revealed that the microspores of all four genotypes remained uninuclear. This developmental stage was found to be the most responsive to embryogenesis induction in many crop plants (<u>Soriano et al., 2013</u>). Of the factors tested, the most crucial for further development of the microspore was the induction medium formulation. Using a relatively complex medium, a few microspores responded (0.05–0.5%) and developed further, while the remainder died within 5–10 days. Microspore derived embryos initiated by a series of random divisions within the surrounding exine wall. The nucleus of uninucleate microspores condensed and reduced in size during the first 2 days in culture. They then divided symmetrically within the first 5–8 days, forming two equal-sized nuclei. This developmental stage is considered the initial stage that is often referred to as sporophytic growth (<u>Soriano et al., 2013</u>). Within another 3–5 days, the nuclei underwent a series of divisions resulting in the formation of multinucleate structures. By approximately the third week of culture, globular stage embryos. To date, growth has



not progressed past this stage of embryo development. Current experiments including adjustment of the osmotic and removal of secondary metabolites which could inhibit (microspore-derived) embryo development are running.

**Figure 9:** Developmental pathways observed in *Cannabis* (industrial Cannabis) microspore culture. (A–C) Male gametophyte development in *Cannabis* during *in vitro* culture. (A) Uninucleate microspores; (B) uninucleate microspores after 3 days in culture media; (C) symmetrically divided microspore with two equally sized nuclei; (D) multinucleate structure without organization and still enclosed in exine; (E) globular multicellular structure with developing exine; and (F) heart-shape embryo with two distinct domains. The nuclei in (A–C) are stained with the nuclear dye 4',6-diamidino-2-phenylindole (DAPI) to indicate viability.



#### In vitro Mutagenesis

A mutation occurs in DNA, naturally or it can also be induced artificially. Most of the genetic variation existing in a gene pool has occurred naturally. These genetic variations can be recombined through conventional breeding practices to develop a novel variety with desired gene traits. Although these spontaneous mutations are frequent, the desired mutation in the desired gene segment altering its biological role is extremely rare. Therefore, mutation induction tools are used in the rapid development of genetic variability in crops. For the last few decades, there were several scientific reports published assessing the impact of an induced mutation in the improvement of crops (Brock, 1971; Broertjes and Van Harten, 1988; Micke, 1999; Oladosu et al., 2016). However, in cannabis research and development is rapidly flourishing, but there are only a few reports on targeted mutation through genetic transformation (Feeney and Punja, 2003; Slusarkiewicz-Jarzina et al., 2005; Sirkowski, 2012; Wahby et al., 2013) and there is no mutant variety introduced at the commercial level. *In vitro* culture techniques, coupled with mutagenesis, has simplified the crop improvement work for both seeds and vegetatively propagated plants (Hussain et al., 2012).

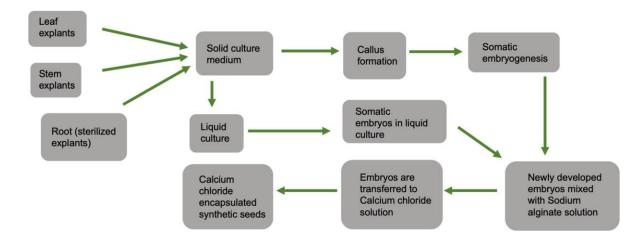


Little efforts have been made and published to establish DH production in cannabis, but once streamlined will open up exciting opportunities for DH mutagenesis as it has been successfully employed in canola (<u>Szarejko, 2003</u>).

#### Synthetic Seed Technology

Synthetic seeds usually refer to artificially encapsulated somatic embryos (<u>Murashige, 1977</u>) but have also been used in reference to encapsulated vegetative tissues that have the potential to develop into a whole plant (auxiliary buds, cell aggregates, shoot buds). Somatic embryos provide the ideal approach to developing synthetic seeds as they often have the ability to survive desiccation and can be treated in much the same way as true seeds. At the same time, other tissues lack this capacity and are less useful (<u>Rihan et al., 2017</u>). As shown in synthetic seeds can be successfully developed by using various explants, media, and encapsulation protocols (<u>Bapat et al., 1987</u>; <u>Corrie and Tandon, 1993</u>; <u>Nyende et al., 2003</u>; <u>Chand and Singh, 2004</u>; <u>Rai et al., 2008</u>; <u>Lata et al., 2009a</u>).

Figure 10: General schematic diagram showing steps for calcium chloride encapsulated synthetic seed production.



Cannabis is generally a cross-pollinating crop, and due to its allogamous nature, it is difficult to maintain existing elite varieties by seed. Typically, a minimum isolation distance of 5 km between breeding nurseries and Cannabis production fields is required to minimize the occurrence of nuisance pollen. Such separation is often difficult to achieve in areas with high Cannabis production intensity. Therefore, *in vitro* propagation using synthetic seed technology is an alternative method for large-scale clonal propagation and germplasm preservation. As the cannabis industry grows, this method may be cheaper and faster than traditional tissue culture methods. Along with the preservation of genetic uniformity, clones produced through this technique are pathogen-free, easy to handle, and transport.

Moreover, in other species, this approach has resulted in increased quality of planting material (<u>Rihan et al., 2017</u>). While cannabis tissue culture methods are still being optimized, <u>Lata et al. (2009a)</u> developed a high-frequency



propagation of axillary buds of *Cannabis* encapsulated in calcium alginate gel. Calcium alginate is a hydrogel that contains nutrients, growth regulators, and sometimes antibiotics.

When directly sown on a substrate, encapsulation aids in the physical protection and establishment and growth of the explant. According to Lata et al. (2009a), gel capsule consisted of 5% sodium alginate with 50 mM CaCl<sub>2</sub>.2H<sub>2</sub>O, and full-strength MS medium supplied with 0.5  $\mu$ M TDZ, and 0.075% plant preservative mixture (PPM). The optimal regrowth and conversion were achieved in MS medium supplemented with antimicrobial components, PPM (0.075%) and TDZ (0.5  $\mu$ M) under *in vitro* conditions. Under *in vivo* condition, the optimal conversion and regrowth were exhibited on 1:1 potting mix-fertilome with coco natural growth medium supplied with MS medium containing 3% sucrose, 0.5% PPM. Clones regenerated from the explants were successfully hardened and transferred to the soil (Lata et al., 2009a).

Another hurdle to *in vitro* propagation is transporting requested strains from the tissue culture facility to the growers in a timely manner. These transportation issues become incredibly challenging for maintaining crop schedules because cannabis crops can take more than 2 months to reach hardening stages, then spend 4 weeks in vegetative growth, then 7 or 8 weeks in flower. Greenhouse or indoor growers require a consistent supply demand to receive a high volume of plantlets every week to start over a new grow room at a very tight on-time delivery schedule, which is the most important metric in their operations. An established and cost-effective synthetic seed encapsulation technique would provide an opportunity to eliminate the transportation challenge.

#### Cryopreservation

Cryopreservation refers to the storage of diverse living materials at below  $-130^{\circ}$ C (Engelmann, 2004). It serves as an alternative conservation approach to the conventional field and *in vitro* (i.e., slow growth) germplasm conservation and is cost-effective over extended periods with minimal space and routine maintenance requirements (Pence, 2011; Engelmann, 2014; Popova et al., 2015). It also assists current breeding programs by providing long-term storage and an easy long-distance exchange of genetic materials (e.g., pollen and meristematic apices and buds). Cryopreservation has been implemented for various plant species using different methods, the most popular and widely applicable, including controlled freezing, vitrification, encapsulation-dehydration, encapsulation-vitrification, and droplet-vitrification (Sakai and Engelmann, 2007; Popova et al., 2015). These methods follow distinct approaches to dehydrate cryopreserving living materials by converting liquid water to a glassy state to avoid the lethal formation of intracellular ice. The selection of methods and the scales of conservation using this approach are strongly determined by genotypes and tissue materials used, which contain different responses to pre- and post-cryopreservation treatments.

Conventional and *in vitro* conservation of cannabis require considerable amounts of space and routine maintenance, have genetic mutations accumulate in the plants. Conventional conservation may expose plants to virulence pathogens. The plants may eventually become susceptible to diseases. The application of cryopreservation can serve as an essential tool for the conservation of various valuable *Cannabis* genotypes with unique attributes and trading the genotypes nationally and internationally in sterile conditions. The first study on applying cryopreservation techniques in *Cannabis* was reported in 1989 using cell suspension cultures (Jekkel et al., 1989). The suspension



cultures were preserved using 10% dimethyl sulfoxide (DMSO) cryoprotectant and a controlled cooling rate of 2°C/min and transfer temperature of  $-10^{\circ}$ C, with a 58% survival rate after cryopreservation of the cultures. A cryopreservation protocol for *Cannabis* shoot tips was recently developed using a droplet-vitrification in liquid nitrogen for long-term conservation of this crop (Uchendu et al., 2019). The report showed that vitrified shoot tips using a cryoprotectant solution of 30% glycerol, 15% ethylene glycol, 15% DMSO in liquid MS medium with 0.4 M sucrose, pH 5.8 had 63% re-growth efficiency. Despite the promising progress made, more studies need to be done on selecting appropriate cryopreservation methods with respect to the tissue types and genotypes, increasing re-growth and survival efficiency of preserved samples, and genetic stability of regenerated plants after using different cryopreservation tools, among others.

## **Germplasm Maintenance**

The *in vitro* condition also raises some issues for concern, primarily when the material is maintained over a long period of time.

# **Clonal Stability** *in vitro* **Culture**

*In vitro* mass-propagation and maintenance of elite germplasm requires genetically stable true-to-type clones. Several factors, such as the number of subcultures, changes in the relationship of auxin/cytokinin, explant type, and a high concentration of growth regulators, may influence the genetic stability of a clone under *in vitro* conditions (Joyce et al., 2003; Sato et al., 2011; Smulders and de Klerk, 2011; Nwauzoma and Jaja, 2013). While carefully selecting explant types and optimizing the conditions above, but depending on the plant species, clonal stability can be obtained during *in vitro* mass-propagation and germplasm conservation of the desired elite genotypes maintained. To date, *Cannabis* plants regenerated from nodal culture, and *in vitro* conserved synthetic seeds ('Encapsulated' nodal segments) have shown no evidence of genetic mutations; however, this has only been evaluated using low numbers of markers (Lata et al., 2010a, 2011). Despite optimizing and using properly *in vitro* conditions that limit somaclonal variations, assessment of clonal stability is required to ensure the regenerated clones are the true-to-type of the donor plants.

# **Somaclonal Variation**

Although clonal propagation and maintenance of elite germplasm require a substantial genetic uniformity among *in vitro* regenerated plantlets, there may be a large possibility of genetic variations, called "somaclonal variation" among these plants and/or relative to the donor plants. Somaclonal variation is commonly a result of genetic alterations and changes in the new *in vitro* plants' epigenetics compared to the original source plants (<u>Miguel and Marum,</u> 2011; <u>Abreu et al., 2014</u>). The frequency and nature of somaclonal variation *in vitro* culture can be influenced by different factors, such as explant source, genotype, *in vitro* techniques, *in vitro* growth conditions, length of the culture period, and the number of subcultures. The use of *de novo* regeneration from highly differentiated tissues



(i.e., roots, leaves, stems, hypocotyls, cotyledons, etc.) is generally considered to produce more somaclonal variation compared to explants with developed meristems (i.e., axillary buds and shoot tips) (Pijut et al., 2012). Most of these factors generate oxidative stress during culture initiation and subsequent subculturing. The explants and the subsequent regenerated plants exposed to the stress may retain genetic changes. For example, protoplast and callus based plant regeneration impose a high degree of oxidative stress; thus, the stress promotes a high mutation rate, whereas plants regenerated through auxiliary branching (e.g., nodes, shoot tip) experience very low oxidative stress, normally resulting in no genetic variation (Zayova et al., 2010; Smulders and de Klerk, 2011; Krishna et al., 2016). Genetic variation can also arise from somatic mutations already present in the explants collected from the donor plant (Karp, 1994). In vitro regeneration of plants can also be genotype-specific, in which genotypes have different degrees of mutation risks and thus strongly determine the formation of somaclonal variation (Alizadeh et al., 2010; Eftekhari et al., 2012; Nwauzoma and Jaja, 2013). The genetic alterations strongly depend on the in vitro techniques used to regenerate in vitro plants. Additionally, despite differences across plant species, cultures maintained for a long period tend to generate high somaclonal variation, and vice versa (Farahani et al., 2011; Jevremovic et al., 2012; Sun et al., 2013). When cultures are getting old and continuously subcultured, the chance of generating genetically less uniform plants is increased (Zayova et al., 2010), but depends upon the plant species. For example, any more than eight subculture cycles increased somaclonal variation in banana (Khan et al., 2011), whereas over 30 subcultures did not cause any detectable somaclonal variations in Cannabis (Lata et al., 2010a).

Although the molecular mechanism of how somaclonal variations generated from a single plant genotype under the same *in vitro* conditions is not fully explored, several potential mechanisms causing genetic alternations and epigenetics have been proposed in different plant species. These mechanisms include changes in chromosome number, point mutations, somatic crossing over and sister chromatid exchange, chromosome breakage and rearrangement, somatic gene rearrangement, DNA replication, changes in organelle DNA, insertion or excision of transposable elements, segregation of pre-existing chimeral tissues, DNA methylation, epigenetic variation, and histone modifications and RNA interference (Sato et al., 2011; Krishna et al., 2016; references therein).

The occurrence of somaclonal variations in regenerated *in vitro* plants may be advantageous or disadvantageous, depending on *in vitro* propagation goals. If *in vitro* propagation aims to generate new variants, obtaining variations among *in vitro* plants can be advantageous that increases genetic diversity for a genotype used. It provides an alternative tool to the breeders for obtaining genetic variability in different plant species, which are either difficult to breed or have narrow genetic bases. On the flip side, when *in vitro* propagation targets to produce multiple true-to-type *in vitro* plants and maintain elite germplasm, the occurrence of subtle somaclonal variations is a severe problem.

#### **Phytocannabinoid Synthesis in the Cannabis Species**

Nature has deftly adorned cannabis species with a spectrum of phytocannabinoids or monoterpenoids that are chemically designed with para-oriented isoprenyl and aralkyl groups (<u>Hanus et al., 2016</u>). Since the discovery of tetrahydrocannabinol (THC) and cannabidiol (CBD) in the early 1960s, there are over 120 cannabinoids that has been reported, and the biosynthesis pathway of these compounds has been greatly improved (<u>Taura et al., 2014</u>); <u>Sirikantaramas et al., 2004</u>; <u>Taura et al., 2007b</u>, 2009; <u>Gagne et al., 2012</u>; <u>Stout et al., 2012</u>; <u>Laverty et al., 2019</u>).



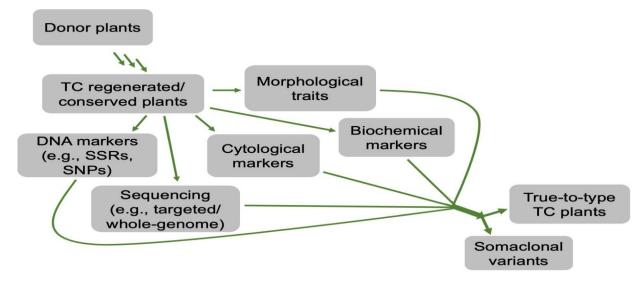
Presumably, cannabigerolic acid (CBGA), the product formed by the alkylation of geranyl diphosphate and olivetol, is the key precursor compound in the synthesis of cannabinoids (Fellermeier and Zenk, 1998). The cyclization event of phenyl components of CBGA, catalyzed by three enzymes – tetrahydrocannabinolic acid synthase (THCAS) (gene bank accession: AB057805), cannabidiolic acid synthase (CBDAS) (gene bank accession: AB292682), and cannabichromenic acid synthase (CBCAS), lead to the formation of three major cannabinoids, THCA, CBDA, and CBCA, respectively (Sirikantaramas et al., 2004; Taura et al., 2007a). Biochemical characterization of the enzymes, THCAS and CBDAS, have demonstrated that the enzymes follow a similar reaction mechanism. In the presence of molecular oxygen, the enzymes use flavin adenine dinucleotide (FAD) cofactor to catalyze CBGA forming THCA and CBDA, and hydrogen peroxide as its chemical biproduct (Sirikantaramas et al., 2004; Taura et al., 2007; Taura et al., 2007b). Although it is a bit unclear, the chemical reaction for CBCAS also believed to use FAD as cofactor and molecular oxygen to complete the enzymatic activity on CBGA. The genes that encode for CBCAS and THCAS are highly similar in the nucleotide level, indicating that CBCAS is also flavoproteins, like the other two enzymes, requiring oxygen to catalyze CBGA to CBCA (Laverty et al., 2019). THCA, CBDA, and CBCA are the major cannabinoids in acidic forms that are synthesized in cannabis plant; upon decarboxylation, these compounds convert into neutral forms, THC, CBD, and CBC respectively (Wang et al., 2016).

## **Determination of Genetic Fidelity**

Variations between regenerated and donor plants can be exhibited at phenotypic, cytological, biochemical, and genetic/epigenetic levels (Hillig, 2005; Miguel and Marum, 2011; Smulders and de Klerk, 2011; Abreu et al., 2014). These variations can be determined through different approaches, such as morphological, cytological, biochemical, and molecular analyses. For morphological traits, changes are not always observed at early developmental stages or may not entirely display the variations. By contrast, the use of cytological and molecular detection approaches determines differences at chromosomal and DNA levels, respectively, regardless of the developmental stages in various plant species (Clarindo et al., 2012; Pathak and Dhawan, 2012; Currais et al., 2013; Abreu et al., 2014; Bello-Bello et al., 2014). To date, several studies have been reported on the use of different molecular markers in *Cannabis* spp. genetic diversity, fingerprinting, etc. These markers include random amplified polymorphic DNA (RAPD), restriction fragment length polymorphisms (RFLP), amplified fragment length polymorphism (AFLP), microsatellites, inter simple sequence repeat (ISSR), short tandem repeat (STR) multiplex, and single nucleotide polymorphisms (SNPs) and PCR Allele Competitive Extension (PACE) assay (Faeti et al., 1996; Kojoma et al., 2002; Alghanim and Almirall, 2003; Gilmore and Peakall, 2003; Hakki et al., 2003; Datwyler and Weiblen, 2006; Mendoza et al., 2009; Lata et al., 2010a; Gao et al., 2014; Dufresnes et al., 2017; Henry et al., 2018). These molecular markers coupled with cytological and morphological analyses (Abreu et al., 2014) are valuable techniques to ensure the genetic stability of in vitro regenerated plants or in vitro conserved germplasm of Cannabis. To date, only ISSR markers have been used to confirm the genetic stability of *Cannabis* synthetic seeds during in vitro multiplication and storage for 6 months under different growth conditions, and in vitro propagated plants over 30 nodal subcultures in culture and hardening in soil for 8 months, compared to the corresponding donor plants (Lata et al., 2010a, 2011).



**Figure 11:** A flow chart depicting different approaches that can be used to determine the genetic stability of *in vitro* regenerated or conserved cannabis plants, compared to its donor counterparts.



## **Projected Contribution of Tissue Culture in the Global Cannabis Industry**

The present global cannabis market is worth \$340 B<sup>2</sup>. To supply cannabis (medical and recreational) to global consumers, a stable supply chain of quality production and value-added product development still needs to be established. Considering the average annual weighted usage base of 110 g per customer (Canaccord Genuity), the global cannabis demand currently could be around 19-20 M kg per year. Major cannabis consumers are in Europe, North America, South America, Asia, and Oceanic parts of the world, with an estimate of 263 million people using the drug in the previous year (European Consumer Stables Report, 2018; World Drug Report, 2019). To produce 20 M kg of cannabis every year, considering a 40-gm yield per plant, would require 500 M clones/seeds a year. An average price of \$10, as, then, the overall present global expected market size for tissue culture clones/manual clones could

be predicted around \$5B. With intensive indoor cultivation, tissue culture clonal planting material can also reduce the risk of fungal and viral diseases, substantially reducing production cost to under \$0.5 per gram to maintain a profitable cannabis production. Considering these global demand scenarios, the supply of clean cannabis clones (pest free, and true to type tested) is an important supply chain component essential for the success and future growth of cannabis industry. To sustain and support the industry growth and make the production cost-effective, optimization in the cannabis tissue culture technology is vital.



 Table 5: Comparison between tissue culture cloning and manual cloning in cannabis.

| Parameter  | Manual Cloning   | Tissue culture cloning           |
|--|--|----------------------------------|
| Space to produce 1000 cuttings (square meters)                     | 3–5  | 0.36                             |
| Clones processed per person per day (count)                        | 200–250  | 1500-2000                        |
| Multiplication Ratio per month                                     | 1–2  | 4–5                              |
| Cost of Production (\$)  | \$3-4  | \$0.5-1                          |
| Clone multiplication in a 3-month cycle                            | 50-80  | 200–250                          |
| Cleanliness  | Chances of contamination   | Disease, pest, and virus free    |
| Vigor  | Chances of reduced vigor from stressed or infected mother plants | Vigor from meristematic reviving |
| Estimated clone production per 10,000 square feet per year (count) | 200,000  | 2,000,000                        |
| Estimated revenue at \$10 per clone                                | \$ 2M  | \$ 20 M                          |

The *in vitro* propagation of cannabis is superior to conventional methods because of disease-free elite plants' production and a high multiplication rate. The cannabis industry is keen to invest in *in vitro* propagation due to (i) saving footprint/production area by shifting a mother room to a tissue culture lab that will be almost 10% the size of the space needed same number of clones.

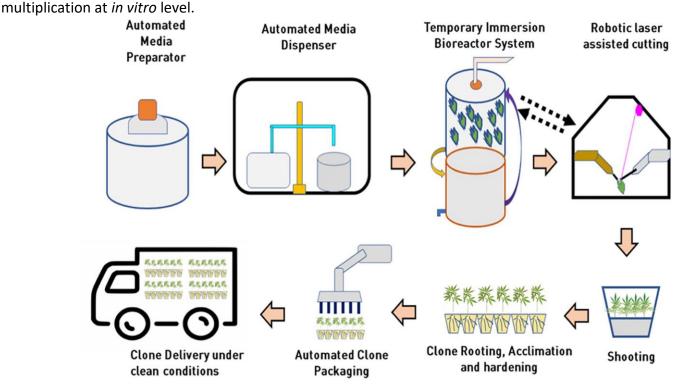
The main hurdle of *in vitro* propagation is the capital cost for the tissue culture lab setup. Setting up a massive largescale production facility can involve a multimillion-dollar investment. Industry and technology will need to continue to improve and reduce costs so that *in vitro* propagation can be affordable for all growers.

In other plants, under a laminar flow hood setting, on an average of 100 plants per hour with 2000 working hours, 200,000 plants can be produced in a year. With an hourly labor cost of \$35 per hour will cost around \$0.35 per tissue culture plant (<u>Sluis, 2005</u>). This is around 60% of the production cost, adding another \$0.15 for other costs (including electricity, resources, and marketing) makes it a baseline cost of \$0.50 per plant. Scale also makes some impact on the cost of production being larger facilities can reduce the cost per plant significantly. These production costs can be as low as \$0.15 per plant if the plants are produced in India, Singapore, China, or Africa where labor costs are comparatively low.

A few biotech companies recently added robotic sub-culturing technology for their cannabis plantlets and developed a fully automated micropropagation system to reduce large-scale operation costs. However, the capital investment to purchase this kind of robotic system is incredibly high currently. Automated technologies for media preparation and dispensing, photoautotrophic bioreactor systems, robotic explant handling, and cutting, transfer laser dissected explants into fresh culture media, and automated acclimatized and hardened plant packaging in future will make cannabis tissue culture industry high throughput and extremely cost-effective for assured "*Just In Time*" supply of pest free, true-to-type cannabis clones. A conceptual model for high throughput automated cannabis *in vitro* clonal mass propagation is depicted in. Robotics has the potential to bring tissue culture cost down by 25% (as low as \$0.15 per plant to compete with low-cost production in some parts of the world). Tissue culture automation technology is slowly progressing, and it will not only bring high-level consistent output but also reduce the cost of production as low as 20 cents per plant.



Figure 12: Integration of automation and bioreactor technologies for mass propagation in cannabis for low cost clonal



The body of Cannabis micropropagation and regeneration literature is poised to undergo substantial growth as regulations around the globe begin to relax. While several existing publications report high rates of MSCs, there have been challenges in replicating the results of these studies across genotypes and research groups. Reports of de novo regeneration are even more limited; their success has been mixed and positive outcomes have been difficult to replicate. These challenges are highlighted by the fact that there are no published reports of regeneration of transgenic plants obtained using traditional molecular and genome editing approaches. In drug-type Cannabis, micropropagation and regeneration protocols suffer from low multiplication rates, poor replicability, and a vast array of starting tissues to choose from, coupled with high diversity in genotypic responses and underwhelming robustness resulting from protocols conceived using single genotypes. Precise methods using multiple genotypes are necessary to develop protocols that can be reliably replicated by other research groups, and innovative new approaches to Cannabis micropropagation are required if developments in Cannabis tissue culture and plant biotechnologies are to keep pace with the needs of the producers and consumers in this burgeoning industry.

The process of developing new varieties through conventional breeding can take 7–12 years, depending on crop species. The progress of cannabis breeding programs is limited due to the difficulty in maintaining selected high-yielding cross-pollinated elite genotypes under field or greenhouse conditions. Therefore, tissue culture techniques are advantageous for cannabis improvement because they can facilitate high multiplication rate and production of disease-free elite plants by overcoming the problems of heterozygosity from cross-pollination. The development of new industrial Cannabis and medical cannabis cultivars with improved traits could be further advanced using genome



editing and other precision breeding tools combined with *in vitro* techniques for regeneration. Unfortunately, Cannabis and cannabis plants' dioecious nature complicates the efforts toward the improvement of specific traits, such as resistance to pests and diseases. Therefore, with the recent legalization, calls for serious targeted efforts are required to advance the regeneration and transformation protocols aiming to enhance the quality and safety of the plants and end products.



https://flow.page/horizonholdings





#### **ABOUT THE TEAM**

Together we (Marc George Ph.D., J Gold Genetics, and Bill Carrington Ph.D.) have formed a compliance Holding company <u>https://flow.page/horizonholdings</u> for the Legal Cannabis/Cannabis and Mushroom Industry Holding subsidiary companies, that are providing Cell to Tissue Culture Media (for Cannabis/Cannabis, Mushrooms, Cactus), Genetics Restructure (breeding cell-seed-tissue, storing, sales), Culturing for sale and R&D (cannabis/Cannabis, mushrooms, and cactus), Cultivation (indoor, field, greenhouse), Fertilizer 17 product line, Education online course accredited University, Extraction (compounds from mushrooms, cactus, cannabis, Cannabis, and hash)(from C02, cold & heat-press, and jar tech), Testing (R&D, molds, DNA, terps, cannabinoids, and compounds), Retail (store and E-Comm), Testing (R&D, molds, DNA, terps, cannabinoids), Retail (store and E-Comm).

Mr. Gold is an executive with 30 years of national and international experience in the cannabis, biotechnology, breeding, and product design industries. Mr. Gold has led the creation, development, and growth of several companies throughout his career. An articulate and trusted voice in the world's cannabis space, Mr. Gold's life-long commitment to our consulting team has accomplished several pioneering Technologies. Created over 1000 genetics for the cannabis industry along with the first CBD strains to Market. Mr. Golds' genetics have been utilized in hundreds of cup-winning events.

Schwegman Lundberg & Woessner, P.A. Alternative to Plant Characteristics without Traditional Genetic Engineering Trademark /patent issues (SLW: 4427.002PRV) https://www.ecronicon.com/ecag/volume5-issue10.php

Plant Characteristics: Traditional Genetic Engineering This invention involves improved methods of generating hybrid plant cells, and hybrid plants, by somatic cell fusion without electric shock. The methods do not require recombinant alteration of cellular chromosomes by currently available genetic engineering procedures. For example, the inventive methods do not involve the transformation of cells by insertion into plant chromosomes or transient expression of coding regions from expression cassettes, expression vectors, viral vectors, plasmids, or other vectors commonly used for genetic engineering. Instead, the nuclei of fused somatic cells can naturally exchange genetic information by homologous recombination using processes like those that occur naturally during the sexual reproduction of plants. New types of hybrid cells are therefore formed that have desirable traits and improved characteristics.

(WO2017007833) HEALTHFUL SUPPLEMENTS, and (WO2018160702) HEALTHFUL SUPPLEMENT FOOD & Schwegman Lundberg & Woessner, P.A (WO2017007833) HEALTHFUL SUPPLEMENTS <u>https://patentscope.wipo.int/search/en/detail.jsf?docId=WO2017007833&recNum=1&maxRec=17&office=&prevFilte</u> <u>r=&sortOption=Pub+Date+Desc&queryString=FP%3A%28healthful+supplements%29&tab=PCT+Biblio</u>

(WO2018160702) HEALTHFUL SUPPLEMENT FOOD that can be given in a solid and or liquid form, and or through a feeding tube. <u>https://patentscope.wipo.int/search/en/detail.jsf?docId=WO2018160702</u>

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