

# High-frequency plant regeneration and genetic fidelity assessment of regenerants by molecular and biochemical markers in Woodland Sage (*Salvia nemorosa* L.)

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## ABSTRACT

*Salvia nemorosa* L. is a valuable medicinal and ornamental plant from Lamiaceae. A high-frequency protocol for in vitro regeneration from *S. nemorosa* leaf explants was developed based on our results. Various plant growth regulators were evaluated for their effects on direct and indirect organogenesis, callusogenesis, shoot multiplication, and rooting of explants at different concentrations. The maximum percentage of direct shoot regeneration (97.5%) and the number of shoots per explant (8.6) were obtained on MS medium supplement with 8  $\mu$ M TDZ plus 8  $\mu$ M NAA. The result revealed that media amended by 16  $\mu$ M 2, 4-D with 8 or 16  $\mu$ M BA was the best treatment for callus induction. Media containing the combination of 8  $\mu$ M TDZ plus 2  $\mu$ M NAA provided the highest rate of shoot induction (84%) and the highest number of shoots (5.4) from calli. Regenerated shoots were multiplied efficiently with an average of 7.4 shoots and 32.4 leaves per explant on media containing 12  $\mu$ M TDZ. In the present study, TDZ and BA treatments led to the highest proliferation rate; however, adenine sulfate was preferred for proliferation because it did not produce hyperhydric plants. The genetic fidelity of 19 plantlets (one donor plant, nine direct and nine indirect regenerants) was appraised by 10 ISSR markers. The molecular analysis revealed that direct organogenesis produced true-to-type plantlets and could be employed for clonal multiplication of Woodland Sage. Indirect regenerants showed polymorphic bands (12.5–33.33%), confirming the somaclonal variation occurrence during indirect organogenesis. Also, HPLC analysis of regenerants showed that two indirect regenerants accumulated higher rosmarinic acid content than mother plants, while three of them accumulated lower rosmarinic acid content. As part of this study, molecular and biochemical markers have been employed to investigate somaclonal variation in in vitro regeneration of the *Salvia* genus for the first time, which could be utilized to develop reliable methods for multiplying elite lines and achieving high-yield lines.

## 1. Introduction

*Salvia* is the largest genus of the Lamiaceae family, with over 1000 species distributed worldwide (Marchev et al., 2014). Members of this genus are a rich source of secondary metabolites such as terpenoids, sterols, flavonoids, and phenolic compounds (Wu et al., 2016). Additionally, extracts of Sage species represent antibacterial, antiviral, antifungal, anti-mutagenic, and anti-inflammatory activities (Jan et al., 2015). These properties caused the widespread use of *Salvia* species in the folk and modern medicinal, cosmetics, and pharmaceutical drugs industry and as a food additive (Wu et al., 2016).

*Salvia nemorosa* L. (Woodland Sage) is one of seventeen endemic *Salvia* species in Iran, which grows wild throughout its mountainous regions (Heydari et al., 2020a; Mirza and Sefidkon, 1999). This species is also distributed in central and southeastern Europe, southern Siberia, Caucasia, and Turkey (Kuzma and Wysokińska, 2003; Skała and Wysockińska, 2004). Woodland Sage is a hardy herbaceous perennial plant with high ornamental and medicinal value and uses widely in the garden and landscape design (Kaprinjak and Fari, 2019; Mirza and Sefidkon, 1999). In Turkish folk medicine, *S. nemorosa* leaves are traditionally applied externally to stop bleeding (Skała and Wysokińska, 2004). In Bulgarian traditional medicine, Woodland Sage is used mainly to treat

**Abbreviations:** 2, 4-D, 2, 4-Dichlorophenoxyacetic acid; AdS, Adenine Sulphate; BA, N6-Benzyladenine; CTAB, Cetyltrimethylammonium Bromide; IAA, Indole-3-Acetic Acid; IBA, Indole-3-butyric acid; KIN, Kinetin; mMS, Modified Murashige and Skoog; NAA,  $\alpha$ -Naphthalene acetic acid; RA, Rosmarinic Acid.

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hemorrhages, stomach aches, furuncles, and diarrhea (Bahadori et al., 2017). This plant has been employed in traditional Iranian medicine to reduce pain and inflammation. Aqueous and ethanolic extracts from the aerial parts showed anti-nociceptive activity with an efficacy similar to classical analgesic drugs (Heydari et al., 2020a; Nikolova and Aneva, 2017). Aerial parts of Woodland Sage are a rich source of phenolic compounds. Rosmarinic acid (RA) is the most abundant phenolic compound detected in this plant (Bahadori et al., 2017). Rosmarinic acid is a phenolic molecule with various applications, varying from food preservatives to cosmetics, and is also a molecule of interest to the pharmaceutical industry (Marchev et al., 2021). The RA possesses anti-inflammatory, anti-oxidative, antibacterial, antiviral, anti-hyperglycemic, anticancer, hepatoprotective, analgesic, cardioprotective, and neuroprotective properties. It is used in modern medicine to treat Alzheimer's disease, diabetes, fibrosis, and various types of cancer such as colorectal, breast, pancreatic, and lung cancers (Hitl et al., 2021; Trócsányi et al., 2020). Salvianolic acids, which have been used to treat cardiovascular disorders, fibrosis, and cancer, are another significant phenolic compound isolated from *S. nemorosa* (Heydari et al., 2020b). Other phenolic compounds detected in *S. nemorosa* aerial parts include caffeic acid, chlorogenic acid, ferulic acid, syringic acid, and p-coumaric acid (Bahadori et al., 2017).

An efficient tissue culture protocol is valid when it produces true-to-type plants (Viehmanna et al., 2016). *In-vitro* condition is very stressful and may lead to genetic and chromosomal instabilities, called somaclonal variation. Although this phenomenon is useful for some breeding purposes, it induces variations that may not be desirable when the goal is clonal multiplication (Krishna et al., 2016). Somaclonal variation, on the other hand, has the potential to induce favorable characteristics and increase variation in a population. It has become a helpful tool in plant improvement with regard to obtaining variants with resistance to abiotic stresses, increased yield and quality, novel phenotypic traits, and aesthetic qualities (Pawelkiewicz et al., 2021). Therefore, concerning the desired objectives, it seems necessary to find out whether somaclonal variation occurs during *in vitro* propagation of plants (Wang and Wang, 2012). Recently, polymerase chain reaction (PCR) techniques, particularly molecular markers, have been routinely used to assess genetic fidelity or somaclonal variation in regenerated plants during *in vitro* multiplication (Viehmanna et al., 2016). There are many techniques available, but the inter simple sequence repeat (ISSR) is advantageous because it eliminates many limitations of other techniques, like the high cost of AFLP, the requirement of species-specific primers for SSR, and the low reproducibility of RAPD (Martínez-Estrada et al., 2017; Peng et al., 2015). ISSR markers provide a highly polymorphic, simple, and fast procedure that incorporates most of the advantages of SSRs and AFLP to the universality of RAPD (Ulvrova et al., 2021). It has been utilized in many species (Dharman and Anilkumar, 2021; Pastelin Solano et al., 2019; Thakur et al., 2021; Ulvrova et al., 2021) to examine somaclonal variation among *in vitro* propagated plants, indicating that it is appropriate for validating genetic integrity after micropropagation (Pawelkiewicz et al., 2021).

*S. nemorosa* is currently "underappreciated," but it could be a promising plant species in the future due to its high ornamental and medicinal values (Kaprinjak and Fari, 2019). Despite its valuable characteristics, it has received limited efforts to improve its growth, biochemical, and ornamental qualities (Heydari et al., 2020b). Plant breeders must compensate for these deficiencies by employing faster biotechnology methods such as somaclonal variation, gene cloning, metabolic pathway engineering, artificial polyploidy, etc. They lead to the generation of plants that vary in traits in a significantly shorter timeframe and with a reduced workload compared to conventional breeding methods, which take 10–15 years and are laborious (Niazian, 2019; Pawelkiewicz et al., 2021). The tissue culture platform is the basis for most of the new biotechnological techniques employed in modern plant breeding. Therefore, establishing a reliable and efficient protocol for *in vitro* regeneration and micropropagation is necessary (Arora et al., 2022).

Though the somaclonal variation in *in vitro* medicinal plant cultured is not well documented, one shall speculate that it is presumably involved in the regulating network of secondary metabolites production. Thus, its possible implication in triggering secondary metabolites production needs to be elucidated (Ngezahayo, 2018). According to our literature review, only one research by Skała and Wysokińska (2004) has been reported on the micropropagation of Woodland Sage. Only a few types and concentrations of plant growth regulators, particularly auxins, were examined in their study, and the genetic integrity of regenerated plants was not reported. Consequently, further studies must be conducted on the *in vitro* propagation of this plant, especially Iranian indigenous genotypes. Our study aimed to develop a reliable and efficient regeneration protocol through direct and indirect organogenesis. Then, *in-vitro* regenerants of *S. nemorosa* were subjected to ISSR marker analyses to assess genetic identity and investigate the occurrence probability of somaclonal variation. Finally, the amount of rosmarinic acid and Phenylalanine ammonia-lyase (PAL) enzyme activity were examined as biochemical markers to evaluate the occurrence of somaclonal variation in regenerated plants.

For the first time, molecular and biochemical markers have been employed in the present study to investigate somaclonal variations in *Salvia* species *in vitro* regeneration. This study is the first report on the validation of genetic fidelity in *S. nemorosa* regeneration. It could be used for large-scale multiplication of this species, which has excellent ornamental and medicinal potential.

## 2. Material and method

### 2.1. Plant material

Seeds of the Woodland Sage were gathered from Khansar, Isfahan province, which is located at 33° 10' 00" N, 50° 23' 00" W. The seeds have been validated in the Herbarium of Research Institute of Forests and Rangelands, Tehran, Iran. They were subjected to a surface sterilization protocol of 70% (v/v) ethanol for 1 min and 2.5% chlorine bleach (NaClO) containing a few drops of Tween 20 for 15 min. After that, rinsed at 3, 5, and 15 min with sterile double distilled water and cultured in glass jars (200 mL) holding 25 mL Murashige and Skoog (MS) basal media (Murashige and Skoog, 1962) solidified by 5.8 g/l plant agar (Plant agar, Duchefa Biochemie, Netherlands). Then, they were placed under cool-white light illumination with a 16 h photoperiod at 24 ± 1 °C. The seeds began germinating after 10 days of culture, and twenty-eight-day-old seedlings were employed as source explants. All explants employed in this study were taken from a single randomly chosen seedling to standardize the experiment. For this purpose, the leaves of the seedling were excised, and the shoots were transferred to a proliferation media described by Skała and Wysokińska (2004), which was MS medium containing 8.9 μM N6-Benzyladenine (BA) and 2.9 μM Indole-3-acetic acid (IAA). Shoot proliferation on the same medium was done until enough individual explants were obtained. In the final step, before the organization of trials, previously gained plantlets were sub-cultured on a MS medium without PGRs. Twenty-one days after the last subculture, mature leaves were cut and used as explant in direct and indirect regeneration experiments.

### 2.2. Effect of PGRs on callus induction and direct regeneration

The goal of the first experiment was to appraise the effect of various plant growth regulators (PGRs) on callus induction and direct regeneration in the leaves of *S. nemorosa*. For this purpose, 1 × 1 cm pieces of *in-vitro* raised leaves as an explant were transferred to the modified MS medium supplemented with BA and Thidiazuron (TDZ) as cytokinin and α-Naphthalene acetic acid (NAA) and 2,4-Dichlorophenoxyacetic acid (2, 4-D) as auxin. Our earlier tests (data not shown) indicated that a modified MS medium (MS basal salts amended with B5 vitamin mixture) was more suitable for callogenesis and organogenesis in Woodland Sage

than MS medium. Also, 100 mg/l of ascorbic acid was added to all media to prevent medium and explants browning by phenol secretion. Thirty-five PGR combinations included BA or TDZ at 0, 4, and 8  $\mu\text{M}$  along with NAA or 2, 4-D at 0, 4, 8, and 16  $\mu\text{M}$  used in this experiment (Table 1). All cultures were kept in the growth chamber under the dark condition at  $24 \pm 1$  °C for 4 weeks. During this period, cultures were monitored regularly, and the data was recorded. The experiment was done based on a completely randomized design with five replications (glass jars). Eight explants were placed per each replication (glass jars). All PGRs used in this study were purchased from Duchefa Biochemie, Netherlands.

### 2.3. Indirect regeneration

The second experiment was conducted to study the effect of PGRs on shoot regeneration from the callus of *S. nemorosa*. Calli was obtained in the first experiment, sub-cultured three times with 14-day intervals on mMS media supplemented by 16  $\mu\text{M}$  2, 4-D + 8  $\mu\text{M}$  BA for proliferation. Then, 1 g of proliferated calli served as an explant in the current experiment. These explants were placed on mMS medium amended by BA or TDZ at 0, 4, 8, and 16  $\mu\text{M}$  in combination with NAA at 0, 2, and 4  $\mu\text{M}$ . All cultures were placed under the dark condition in the growth chamber at  $24 \pm 1$  °C for 6 weeks. During this period, cultures were monitored regularly, and data was recorded. A completely randomized design with five replications (glass jars) was adopted for this experiment. Five explants were cultured per each replication (glass jars).

### 2.4. Shoot multiplication

The third experiment evaluated the influence of various types and concentrations of cytokinins on shoot multiplication of *S. nemorosa*. Shoots regenerated in previous experiments were cultured on mMS medium without PGRs for 28 days. These shoots were employed as the source of explants in the proliferation trial. At first, the leaves are excised, and then the shoots with a length of approximately 3 cm contain three nodes cultured on mMS medium supplemented by BA, TDZ, Kinetin (KIN), and adenine sulphate(AdS) at 0, 4, 8, and 12  $\mu\text{M}$  with or without 2  $\mu\text{M}$  NAA for 4 weeks. The experiment was done with five replications in a completely randomized design and five explants incubated in each replication. All cultured media were placed under cool-white light illumination with 16 h photoperiod at  $24 \pm 1$  °C. In this experiment, multiplied plantlets with a glassy appearance and brittle, translucent, and wrinkled leaves were characterized as hyperhydric shoots based on Gao et al. (2017).

### 2.5. Rooting and acclimatization

In a fourth experiment, some auxins were administered to regenerated shoots to observe their effects on root formation. 2–3 cm basal end of proliferated and elongated shoots were excised and transferred to root induction media which was constructed of  $\frac{1}{2}$  mMS media supplemented with IAA, NAA, and IBA (Indole-3-butyric acid) at 0, 2, 4, 6, and 8  $\mu\text{M}$  plus 500 mg/l activate charcoal and 100 mg/l ascorbic acid. The experiment was conducted on a completely randomized design with five replications (glass jars). In each replication, 10 explants were cultured.

**Table 1**

Effect of plant growth regulators on organogenesis and callogenesis of *S. nemorosa* from leave explants.

Number	Plant Growth Regulators ( $\mu\text{M}$ )				Callus Initiation (%)	Day to Callus Initiation	Callus Fresh Weight (g)	Regeneration (%)	Day to Regeneration Initiation	Number of Regenerated Shoots
	NAA	2, 4-D	BA	TDZ						
1	0	0	0	0	0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
2			4		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
3			8		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
4				4	0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	37.5 $\pm$ 5.5 <sup>f</sup>	16.2 $\pm$ 0.58 <sup>bc</sup>	2.4 $\pm$ 0.50 <sup>g</sup>
5				8	0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	50 $\pm$ 3.9 <sup>e</sup>	15.2 $\pm$ 0.37 <sup>cd</sup>	4.2 $\pm$ 0.20 <sup>f</sup>
6	4				25.0 $\pm$ 3.9 <sup>g</sup>	15.2 $\pm$ 0.8 <sup>a</sup>	0.11 $\pm$ 0.003 <sup>k</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
7	8				35.0 $\pm$ 4.6 <sup>f</sup>	14.4 $\pm$ 0.5 <sup>ab</sup>	0.13 $\pm$ 0.005 <sup>j</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
8	16				62.5 $\pm$ 3.9 <sup>d</sup>	13.8 $\pm$ 0.7 <sup>b</sup>	0.14 $\pm$ 0.006 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
9	4	4			0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	37.5 $\pm$ 5.5 <sup>f</sup>	18.8 $\pm$ 0.73 <sup>a</sup>	2.4 $\pm$ 0.50 <sup>g</sup>
10	8	4			0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	52.5 $\pm$ 7.2 <sup>e</sup>	18.4 $\pm$ 0.67 <sup>a</sup>	10 $\pm$ 0.44 <sup>g</sup>
11	16	4			65.0 $\pm$ 7.2 <sup>d</sup>	11.2 $\pm$ 0.3 <sup>de</sup>	0.19 $\pm$ 0.003 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
12	4	8			42.5 $\pm$ 3.0 <sup>ef</sup>	13.4 $\pm$ 0.7 <sup>b</sup>	0.15 $\pm$ 0.003 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
13	8	8			0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	52.5 $\pm$ 4.6 <sup>e</sup>	17 $\pm$ 0.70 <sup>b</sup>	5.2 $\pm$ 0.58 <sup>e</sup>
14	16	8			0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	60 $\pm$ 6.1 <sup>de</sup>	14.2 $\pm$ 0.58 <sup>bc</sup>	4.6 $\pm$ 0.50 <sup>ef</sup>
15		4			37.5 $\pm$ 5.5 <sup>f</sup>	13.4 $\pm$ 0.6 <sup>b</sup>	0.13 $\pm$ 0.005 <sup>j</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
16		8			50.0 $\pm$ 3.2 <sup>e</sup>	11.2 $\pm$ 0.6 <sup>de</sup>	0.15 $\pm$ 0.003 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
17		16			77.5 $\pm$ 8.2 <sup>c</sup>	10.6 $\pm$ 0.6 <sup>def</sup>	0.17 $\pm$ 0.007 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
18	4	4			47.5 $\pm$ 2.5 <sup>e</sup>	12.0 $\pm$ 0.7 <sup>cd</sup>	0.14 $\pm$ 0.005 <sup>j</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
19	8	4			85.0 $\pm$ 4.6 <sup>bc</sup>	11.2 $\pm$ 0.3 <sup>de</sup>	0.19 $\pm$ 0.003 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
20	16	4			100 $\pm$ 0.0 <sup>a</sup>	8.8 $\pm$ 0.3 <sup>gh</sup>	0.26 $\pm$ 0.007 <sup>d</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
21	4	8			50.0 $\pm$ 3.9 <sup>e</sup>	11.2 $\pm$ 0.5 <sup>de</sup>	0.17 $\pm$ 0.007 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
22	8	8			87.5 $\pm$ 5.5 <sup>b</sup>	9.8 $\pm$ 0.3 <sup>efg</sup>	0.23 $\pm$ 0.005 <sup>e</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
23	16	8			100 $\pm$ 0.0 <sup>a</sup>	8.4 $\pm$ 0.3 <sup>h</sup>	0.27 $\pm$ 0.014 <sup>c</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
24	4		4		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	65 $\pm$ 6.1 <sup>cd</sup>	14.2 $\pm$ 0.58 <sup>bc</sup>	4 $\pm$ 0.32 <sup>f</sup>
25	8		4		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	72.5 $\pm$ 6.1 <sup>c</sup>	15.6 $\pm$ 0.68 <sup>cd</sup>	6.2 $\pm$ 0.58 <sup>d</sup>
26	16		4		60.0 $\pm$ 4.6 <sup>d</sup>	11.0 $\pm$ 0.2 <sup>de</sup>	0.26 $\pm$ 0.007 <sup>d</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
27	4		8		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	95 $\pm$ 3.0 <sup>a</sup>	13.4 $\pm$ 0.74 <sup>f</sup>	8.0 $\pm$ 0.63 <sup>ab</sup>
28	8		8		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	97.5 $\pm$ 2.5 <sup>a</sup>	13.8 $\pm$ 0.58 <sup>ef</sup>	8.6 $\pm$ 0.67 <sup>a</sup>
29	16		8		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	82.5 $\pm$ 5.1 <sup>b</sup>	14.6 $\pm$ 0.40 <sup>de</sup>	7.4 $\pm$ 0.67 <sup>bc</sup>
30		4		4	50.0 $\pm$ 3.9 <sup>e</sup>	13.2 $\pm$ 0.8 <sup>bc</sup>	0.18 $\pm$ 0.006 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
31		8		4	87.5 $\pm$ 5.5 <sup>b</sup>	10.6 $\pm$ 0.6 <sup>def</sup>	0.21 $\pm$ 0.004 <sup>f</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
32		16		4	100 $\pm$ 0.0 <sup>a</sup>	9.8 $\pm$ 0.6 <sup>efg</sup>	0.29 $\pm$ 0.007 <sup>b</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
33	4		8		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	65 $\pm$ 6.1 <sup>cd</sup>	14.8 $\pm$ 0.37 <sup>de</sup>	6.6 $\pm$ 0.67 <sup>cd</sup>
34	8		8		0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>i</sup>	0.0 $\pm$ 0.0 <sup>l</sup>	60 $\pm$ 6.1 <sup>de</sup>	15.6 $\pm$ 0.67 <sup>cd</sup>	6.2 $\pm$ 0.67 <sup>d</sup>
35	16		8		100 $\pm$ 0.0 <sup>a</sup>	9.4 $\pm$ 0.4 <sup>gh</sup>	0.32 $\pm$ 0.009 <sup>a</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>g</sup>	0.0 $\pm$ 0.0 <sup>h</sup>

Mean  $\pm$  SD values in a column, followed by different letters, were significantly different according to the Duncan's multiple range test ( $P \leq 0.05$ ).

All cultures were incubated at  $24 \pm 1$  °C for a 16 h photoperiod under cool-white light illumination. After 14-days, shoots were taken out of the agar medium, and data were recorded. After four weeks, the well-rooted plantlets (176 plantlets) were excised from the culture medium, and the agar was removed by rinsing with sterile distilled water. They were then transferred into plastic pots including cocopeat, peat moss, and perlite (2:1:1). A half strength of MS basal liquid was used to moisten the soil mixture, and pots were covered with transparent polythene bags to hold high humidity. Then, they were maintained in growth chambers at  $24 \pm 1$  °C under a 16 h light and 8 h dark photoperiod at a photon flux rate of  $60 \mu\text{mol m}^{-2} \text{s}^{-1}$  provided by cool fluorescent lamps. After 10 days the polythene bags were removed gradually from the pots and the acclimatized plantlets were finally transferred to the greenhouse.

### 2.5.1. Phenolic compounds extraction

The leaf and stem tissues were ground into a fine powder after being oven-dried for 48 h at 45 degrees Celsius until constant dry weight. 500 mg of powdered samples were mixed well with 10 cc methanol. The mixture was then sonicated at room temperature for 30 min. The extracts were then filtered using Whatman no. 1 filter paper. A rotary evaporator was used to remove the solvent from the extracts. Then, 10 mg of dried extract was dissolved in 10 mL methanol and kept for further analysis.

### 2.5.2. Quantification of rosmarinic acid

A high-performance liquid chromatography method was used to assess the content of rosmarinic acid (RA), according to Heydari et al. (2020b). The mobile phase consisted of methanol (solvent A) and distilled water (solvent B) with a 1 mL/min flow rate. Gradient elution was carried out in a following system: 0–30 min, 88% A, 12% B; 30–45 min, 80% A, 20% B; 45–60 min, 70% A, 30% B. The oven temperature was maintained at 25 °C, and peaks were monitored at 195–400 nm wavelength. For the calibration curve, 20–100  $\mu\text{L}$  of the methanolic standard solution of rosmarinic acid was applied in triplicate onto the HPLC. The rosmarinic acid calibration curve was depicted by plotting peak area versus concentrations. Using the equation obtained from the calibration curve, the concentration of the unknown samples was determined. The rosmarinic acid chromatographic peak was recognized by comparing its retention time with its reference standards.

### 2.5.3. PAL activity

Phenylalanine ammonia-lyase (PAL) activity was measured as described by Dong et al. (2010). The leaf tissue (2 g) was homogenized with 5 mL of extraction buffer utilizing a mortar and pestle. The homogenate was filtrated through four layers of cheesecloth and centrifuged at 12,000g for 15 min at 4 °C. The supernatant was used as a crude enzyme. The reaction mixture (3 mL) contained 0.5 mL of the crude enzyme, 16 mM of L-phenylalanine, 50 mM of Tris HCl buffer (pH8.9), and 3.6 mM of NaCl. The mixture was incubated at 37 °C for 60 min and the reaction was stopped by 500  $\mu\text{L}$  of 6 M HCl. The reaction mixture was then centrifuged for 10 min at 12,000 $\times$ g. Production of cinnamic acid was measured as an increase in absorbance at 290 nm. The enzyme activity was calculated per unit weight of protein.

## 2.6. DNA isolation and ISSR analysis

For genetic fidelity studies, nine *in-vitro* raised plantlets through indirect regeneration, nine direct regenerated plantlets, and a control plant were chosen randomly. The modified CTAB method described by Aleksic et al. (2012) was used for the extraction of genomic DNA extraction from fresh leaves. The quality of the extracted DNA was certified in 0.8% agarose (Duchefa Biochemie, Netherlands) stained with 10  $\mu\text{L}$  Gel Stain (Invitrogen, USA) and using a Nanodrop Spectrophotometer (Thermo Scientific, USA). All DNA samples were adjusted to the final concentration of 50 ng/ $\mu\text{L}$  for PCR and preserved in the refrigerator at  $-20$  °C. Out of 18 tested primers (Integrated DNA

Technologies, Belgium), 10 produced at least three clear and scorable bands, and these primers were used for further PCR analysis. The reaction mixture sets in the volume of 20  $\mu\text{L}$  included 2.5  $\mu\text{L}$  template DNA (50 ng/ $\mu\text{L}$ ), 1.5  $\mu\text{L}$  primer, 6  $\mu\text{L}$  sterile double-distilled  $\text{H}_2\text{O}$ , and 10  $\mu\text{L}$  2x Master Mix [150 mM Tris–HCl, pH 8.5, 40 mM  $(\text{NH}_4)_2\text{SO}_4$ , 0.2% Tween 20, 3 mM  $\text{MgCl}_2$ , 400  $\mu\text{M}$  of each dNTPs, 100 U/mL Taq-Purple DNA polymerase, Inert red dye and stabilizer (AMPLIQON, Denmark)]. Amplifications were performed in a Q-Cycler Thermal Cycler (Bio-Rad, USA), programmed with an initial denaturation at 94 °C for 5 min, followed by 40 cycles of denaturation at 94 °C for 1 min; annealing temperatures ranged from 47 to 53 °C for 1 min, and extension at 72 °C for 2 min, with the final extension at 72 °C for 8 min. Amplicons were separated on 1.2% agarose gel stained by 10  $\mu\text{L}$  Gel Stain and run at 75 V for about 1.2 h. The amplified stained fragments were visualized on an agarose gel in Gel Documentation System under UV illumination.

## 2.7. Statistical analysis

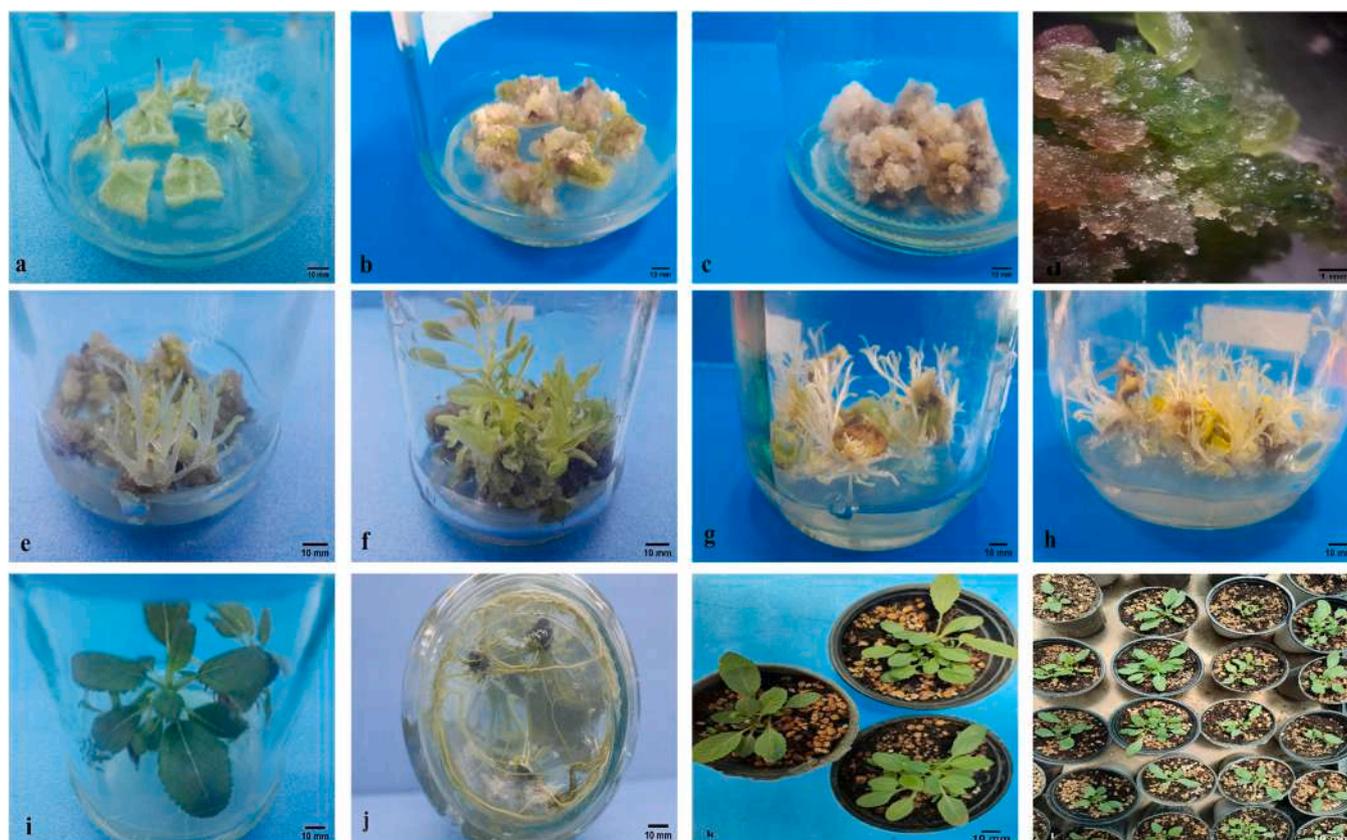
All experiments were done based on a completely randomized design with five replications. The SAS 9.4 software was used for analyzing the data. Duncan's multiple range test (DMRT) was served at the 5% probability level to compare the means. ISSR analysis was done by scoring the reproducible and clear bands based on their presence (1) or absence (0) in the agarose gel.

## 3. Results

### 3.1. Effect of PGRs on callus induction and direct regeneration

The different types and concentrations of PGRs and their combinations used for callus induction and shoot regeneration from leaf explants had significant ( $P \leq 0.05$ ) effects on measured traits (Table 1). No responses were observed from explants incubated on PGRs free medium (control). After about a week of culture, most of the explants began to brown and yellow. However, some of them remained green without any morphogenesis response. It indicated that the presence of an appropriate plant growth regulator in the *S. nemorosa* culture medium was essential for the initiation of organogenesis and callus formation. Among the two applied auxins in this experiment, most treatments containing 2, 4-D resulted in callogenesis (Fig. 1-b). However, PGR combinations containing NAA showed different responses depending on the type and concentration of participant cytokinins. 2, 4-D alone and in combination with BA and TDZ at all levels except two combinations (4  $\mu\text{M}$  2, 4-D + 8  $\mu\text{M}$  TDZ and 8  $\mu\text{M}$  2, 4-D + 8  $\mu\text{M}$  TDZ treatments) resulted in callus induction. On the other hand, regeneration was achieved in media containing NAA except for treatments with a high ratio of NAA to cytokinins, which led to callus induction (16  $\mu\text{M}$  NAA + 4  $\mu\text{M}$  TDZ and 16  $\mu\text{M}$  NAA + 4  $\mu\text{M}$  BA). Out of the 35 combined treatments of PGRs used in these experiments, only 19 treatments led to callogenesis, which significantly ( $P \leq 0.05$ ) differed. A week after culture, in some treatments, including high auxin levels, the explants began to swell and formed a callus on their cut edges (Fig. 1, a-b).

At the end of the culture period, white friable, creamy compact, and white compact calli were developed based on PGRs treatments. Although acceptable callus formation has been attained in the presence of NAA or 2, 4-D (4–16  $\mu\text{M}$ ) in the culture media, their combination with cytokinin (TDZ or BA) accelerated callus initiation and enhanced the percent of callus induction. The result showed that the highest callus induction frequency (100%) was obtained in media amended with 16  $\mu\text{M}$  2, 4-D + 4, and 8  $\mu\text{M}$  BA and TDZ (Table 1). Results also revealed that callus formation and callus weight increased as cytokinin concentration increased. Our results showed that for callus initiation under *in-vitro* conditions, NAA was generally less effective than 2, 4-D. No significant ( $P \leq 0.05$ ) difference was found in the case of different cytokinins (BA and TDZ) effects on morphogenetic response. However, their responses are dependent on the companion auxin. Both caused



**Fig. 1.** - Morphogenesis in *S. nemorosa*. **a.** Callus initiated in cut edges of explants after 7 days from incubation; **b.** Callus formation at the end of experiment; **c.** Callus proliferation; **d.** Shoot primordia formation on calli; **e-f.** Indirect shoot regeneration; **g-h.** Direct shoot regeneration; **i-j.** Rooted plantlets; **k-l.** Acclimatized plant.

callogenesis in most combinations containing 2, 4-D, and induced organogenesis in NAA, including treatments. The results showed that TDZ alone induced adventitious shoots in explants (Fig. 1, g-h). However, BA was unable to develop any morphogenetic response without the presence of auxin. Overall, for direct *in-vitro* regeneration of *S. nemorosa*, TDZ showed more effective than BA. However, media containing BA and TDZ produced the highest number of new plantlets and shoot regeneration frequency, respectively. It's found that the greatest direct regeneration rate was 97.5% and 95% with 8  $\mu\text{M}$  TDZ + 8  $\mu\text{M}$  NAA and 8  $\mu\text{M}$  TDZ + 16  $\mu\text{M}$  NAA treatments, respectively. Moreover, the last PGRs combinations gave the highest shoot number (8.6 and 8 per explant) and the least time (13.8 and 14.6 days) to organogenesis initiation. In fact, the combination of BA with NAA in culture media led to the regeneration of a few number of shoots when compared to media containing TDZ.

### 3.2. Indirect regeneration

The result indicated that media without PGRs failed to regenerate any shoots from the callus, and the callus began to brown and eventually ruined two weeks after incubation. Adventitious shoot initiation started from the explants in other mediums after about 23 days (Fig. 1, d-f). The best results were obtained when callus pieces were placed on media amended with 8  $\mu\text{M}$  TDZ + 2  $\mu\text{M}$  NAA which induced the highest number of shoots (5.8) per explants and regeneration frequency (84%). TDZ produced higher shoots per explant than BA, according to the results (Table 2). It was also found that the result of TDZ treatment was better at lower concentrations and its optimum level was 8  $\mu\text{M}$ . By increasing the TDZ concentration, the number of regenerated shoots decreased. On the other hand, increasing the BA concentration increased the regeneration rate, and its best concentration was 16  $\mu\text{M}$ . The results of this experiment indicate that the presence of low doses of NAA in the regeneration media increased the adventitious shoot production in the

**Table 2**

Effect of plant growth regulators on organogenesis from calli in *S. nemorosa*.

Number	Plant Growth Regulators ( $\mu\text{M}$ )			Regeneration (%)	Number of Regenerated Shoots
	TDZ	BA	NAA		
1	0	0	0	0.0 $\pm$ 0.0 <sup>h</sup>	0.0 $\pm$ 0.0 <sup>h</sup>
2	4	0	0	44 $\pm$ 4.0 <sup>efg</sup>	3.8 $\pm$ 0.58 <sup>c-f</sup>
3	8	0	0	64 $\pm$ 4.0 <sup>bc</sup>	4.6 $\pm$ 0.50 <sup>abc</sup>
4	16	0	0	48 $\pm$ 4.8 <sup>def</sup>	3 $\pm$ 0.31 <sup>efg</sup>
5		4	0	32 $\pm$ 4.8 <sup>g</sup>	2.6 $\pm$ 0.24 <sup>fg</sup>
6		8	0	44 $\pm$ 4.0 <sup>efg</sup>	3.8 $\pm$ 0.58 <sup>c-f</sup>
7		16	0	52 $\pm$ 4.8 <sup>cde</sup>	4.0 $\pm$ 0.31 <sup>b-e</sup>
8	4	2	2	68 $\pm$ 4.8 <sup>b</sup>	4.2 $\pm$ 0.37 <sup>b-e</sup>
9	8	2	2	84 $\pm$ 4.0 <sup>a</sup>	5.6 $\pm$ 0.24 <sup>a</sup>
10	16	2	2	44 $\pm$ 4.0 <sup>efg</sup>	3.6 $\pm$ 0.40 <sup>c-f</sup>
11		4	2	40 $\pm$ 6.3 <sup>efg</sup>	3 $\pm$ 0.40 <sup>efg</sup>
12		8	2	48 $\pm$ 4.8 <sup>def</sup>	4.4 $\pm$ 0.24 <sup>a-d</sup>
13		16	2	60 $\pm$ 6.3 <sup>bcd</sup>	5.2 $\pm$ 0.58 <sup>ab</sup>
14	4	4	4	60 $\pm$ 6.3 <sup>bcd</sup>	3.2 $\pm$ 0.48 <sup>d-g</sup>
15	8	4	4	84 $\pm$ 4.0 <sup>a</sup>	5.2 $\pm$ 0.20 <sup>ab</sup>
16	16	4	4	52 $\pm$ 4.8 <sup>cde</sup>	4.2 $\pm$ 0.58 <sup>b-e</sup>
17		4	4	36 $\pm$ 4.0 <sup>f<sup>g</sup></sup>	2.2 $\pm$ 0.58 <sup>g</sup>
18		8	4	44 $\pm$ 4.0 <sup>efg</sup>	4.4 $\pm$ 0.24 <sup>a-d</sup>
19		16	4	64 $\pm$ 4.0 <sup>bc</sup>	4.6 $\pm$ 0.50 <sup>abc</sup>

Mean  $\pm$  SD values in a column, followed by different letters, were significantly different according to the Duncan's multiple range test ( $P \leq 0.05$ ).

presence of an appropriate ratio of cytokinin to auxin. It is concluded that in our experiment, the best result of PGRs combinations was the ratio of 4-1 cytokinin to auxin.

### 3.3. Shoot multiplication

In this experiment, different concentrations of TDZ, BA, KIN, and AdS (0, 4, 8, and 12  $\mu\text{M}$ ) were evaluated to attain strong and healthy shoots. A significant ( $P \leq 0.05$ ) difference was observed between treatments. Regarding the number of regenerated shoots and leaves per each explant, TDZ showed a better response than others. However, shoot length was significantly ( $P \leq 0.05$ ) more in KIN and AdS supplemented media. New shoots began to form ten days after the explant establishment. After four weeks, media enriched with 12  $\mu\text{M}$  TDZ produced the highest number of adventitious shoots (7.4) and leaves (32.4) per explant (Table 3). However, the high percentage of shoots (54%) grown on that media were shown hyperhydricity symptoms. Raising the concentration of TDZ and BA in the media from 4 to 12  $\mu\text{M}$  increased the multiplication rate. However, the percentage of hyperhydricity (undesirable response) in media containing TDZ raised from 28% to 54% and BA from 22% to 46%. Media supplemented with adenine sulphate (4–12  $\mu\text{M}$ ) produced the highest shoot length (2.17–2.72 cm) without hyperhydricity (Fig. 1-i). Although the proliferation rate in media containing AdS is lower than TDZ and BA treatments, the appropriate shoot length was gained, and hyperhydricity did not occur. Hence, it seems the last one should be media for the multiplication of the Woodland Sage.

### 3.4. Rooting and acclimatization

The *in-vitro* grown shoots were transferred to the half-strength mMS medium containing IAA, NAA, and IBA at 0, 2, 4, 6, and 8  $\mu\text{M}$ . The results revealed that all explants incubated in these treatments produced healthy roots (Fig. 1-j). The highest and significant ( $P \leq 0.05$ ) root number per explants were obtained at 4 and 6  $\mu\text{M}$  NAA (6.4, 6.2 respectively) and 6  $\mu\text{M}$  IAA (6.2) treatments (Table 4). However, NAA at all concentrations provided the highest root number and length rather than IAA. However, media containing NAA (all concentrations) and IBA (only at 6 and 8  $\mu\text{M}$ ) formed a callus. Generally, three evaluated auxins showed a decrement in root number and length at 8  $\mu\text{M}$ . However, it formed thickened roots. Overall, it has been concluded that IAA at the rate of 6  $\mu\text{M}$  was the optimum concentration for the *in-vitro* rooting of *S. nemorosa*. *In vitro*, rooted plantlets were successfully acclimatized in a growth chamber and then transferred to the greenhouse. A total of 176 plantlets were transferred to the greenhouse for acclimatization, and the percentage of surviving plants was 94% after 12 weeks. Acclimatized plants appeared normal and did not exhibit any morphological abnormalities (Fig. 1, k-l).

### 3.5. ISSR analysis

Genetic stability investigation using genomic DNA of the donor and

**Table 3**  
Effect of plant growth regulators on proliferation of nodal explants in *S. nemorosa*.

Number	TDZ	Plant Growth Regulators ( $\mu\text{M}$ )			Mean shoot number per explant	Mean shoot length (cm)	Mean number of leaves per explant	Hyperhydricity frequency (%)
		BA	KIN	AdS				
1	0	0	0	0	1.4 ± 0.24 <sup>g</sup>	1.12 ± 0.06 <sup>i</sup>	6.6 ± 0.67 <sup>g</sup>	0.0 ± 0.0 <sup>f</sup>
2	4				5.0 ± 0.54 <sup>bcd</sup>	1.44 ± 0.04 <sup>h</sup>	23.2 ± 1.15 <sup>cd</sup>	28.0 ± 1.42 <sup>c</sup>
3	8				6.2 ± 0.37 <sup>ab</sup>	1.81 ± 0.14 <sup>fg</sup>	27.6 ± 1.07 ± b	42.0 ± 2.28 <sup>b</sup>
4	12				7.4 ± 0.50 <sup>a</sup>	2.1 ± 0.07 <sup>cde</sup>	32.4 ± 0.92 <sup>a</sup>	54.0 ± 2.28 <sup>a</sup>
5		4			3.2 ± 0.58 <sup>ef</sup>	1.62 ± 0.06 <sup>gh</sup>	15.2 ± 1.24 <sup>f</sup>	22.0 ± 2.28 <sup>d</sup>
6		8			4.4 ± 0.50 <sup>cde</sup>	1.83 ± 0.03 <sup>efg</sup>	22.8 ± 1.06 <sup>cd</sup>	30.0 ± 2.09 <sup>c</sup>
7		12			6.4 ± 0.50 <sup>ab</sup>	2.21 ± 0.06 <sup>cd</sup>	25.4 ± 0.050 <sup>bc</sup>	46.0 ± 3.16 <sup>b</sup>
8			4		2.8 ± 0.37 <sup>efg</sup>	1.94 ± 0.06 <sup>def</sup>	16.8 ± 0.60 <sup>f</sup>	0.0 ± 0.0 <sup>f</sup>
9			8		3.2 ± 0.37 <sup>ef</sup>	2.36 ± 0.05 <sup>bc</sup>	15 ± 0.63 <sup>f</sup>	10.0 ± 1.09 <sup>e</sup>
10			12		5.2 ± 0.80 <sup>bc</sup>	2.81 ± 0.18 <sup>a</sup>	20.6 ± 0.92 <sup>de</sup>	28.0 ± 1.42 <sup>c</sup>
11				4	2.6 ± 0.40 <sup>fg</sup>	2.17 ± 0.09 <sup>cd</sup>	13.4 ± 0.81 <sup>f</sup>	0.0 ± 0.0 <sup>f</sup>
12				8	3.6 ± 0.50 <sup>def</sup>	2.56 ± 0.14 <sup>ab</sup>	15.2 ± 0.80 <sup>f</sup>	0.0 ± 0.0 <sup>f</sup>
13				12	4.2 ± 0.58 <sup>c-e</sup>	2.72 ± 0.05 <sup>a</sup>	18.4 ± 0.81 <sup>e</sup>	0.0 ± 0.0 <sup>f</sup>

Mean  $\pm$  SD values in a column, followed by different letters, were significantly different according to the Duncan's multiple range test ( $P \leq 0.05$ ).

**Table 4**

Effect of Different Auxins on Rooting of explants in *S. nemorosa*.

Number	Plant Growth Regulators ( $\mu\text{M}$ )			Root Number	Root Length (cm)
	NAA	IAA	IBA		
1	0			0.00 ± 0.00 <sup>d</sup>	0.00 ± 0.00 <sup>g</sup>
2	2			5.00 ± 0.37 <sup>b</sup>	3.75 ± 0.14 <sup>ef</sup>
3	4			6.66 ± 0.60 <sup>a</sup>	4.69 ± 0.06 <sup>ab</sup>
4	6			6.50 ± 0.37 <sup>a</sup>	4.92 ± 0.05 <sup>a</sup>
5	8			4.66 ± 0.60 <sup>b</sup>	4.21 ± 0.05 <sup>cd</sup>
6		2		3.50 ± 0.37 <sup>c</sup>	3.57 ± 0.06 <sup>f</sup>
7		4		5.00 ± 0.37 <sup>b</sup>	4.40 ± 0.05 <sup>bc</sup>
8		6		6.50 ± 0.37 <sup>a</sup>	4.76 ± 0.08 <sup>a</sup>
9		8		5.33 ± 0.24 <sup>ab</sup>	4.39 ± 0.05 <sup>bc</sup>
10			2	3.50 ± 0.58 <sup>c</sup>	3.67 ± 0.4 <sup>f</sup>
11			4	5.66 ± 0.60 <sup>ab</sup>	4.43 ± 0.04 <sup>c</sup>
12			6	4.50 ± 0.31 <sup>b</sup>	4.05 ± 0.05 <sup>de</sup>
13			8	6.00 ± 0.31 <sup>a</sup>	3.77 ± 0.08 <sup>ef</sup>

Mean  $\pm$  SD values in a column, followed by different letters, were significantly different according to the Duncan's multiple range test ( $P \leq 0.05$ ).

*in-vitro* raised plantlets from two regeneration methods was carried out to evaluate the possibility of the somaclonal variation using the ISSR marker. Nine randomly selected *in-vitro* propagated plantlets from organogenesis methods (direct and indirect organogenesis) were used and compared with the mother plant. A total of 1240 fragments of between 100 and 1100 bp were amplified using ISSR primers (Table 5). The number of bands per primer differed from four to eight, with an average of 6.4. In the case of the nine directly regenerants and mother plants, the banding patterns generated by ISSR primers are monomorphic, indicating no genetic variation within the samples. However, banding patterns among indirectly regenerated plantlets and donor plants revealed the occurrence of polymorphism. In this case, seven primers of total 10 primers namely UBC-807, UBC-809, UBC-810, UBC-811, UBC-834, UBC-843 and UBC-847 showed polymorphism in 12.5–33.33%. The primers that produced more polymorphic bands (2 bands) were UBC-807 and UBC-810, with 33.33% and 28.57%, respectively (Fig. 2).

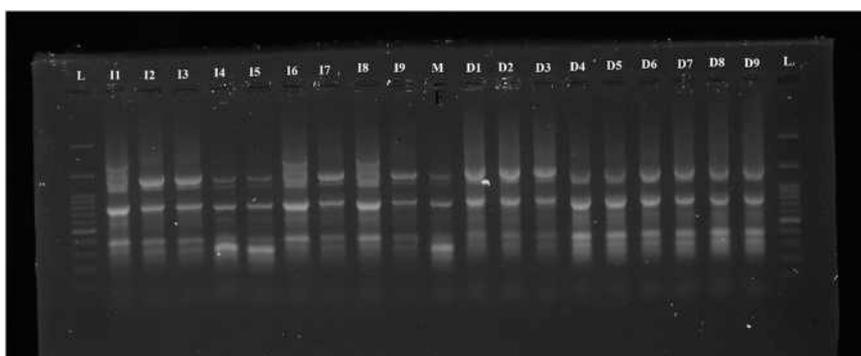
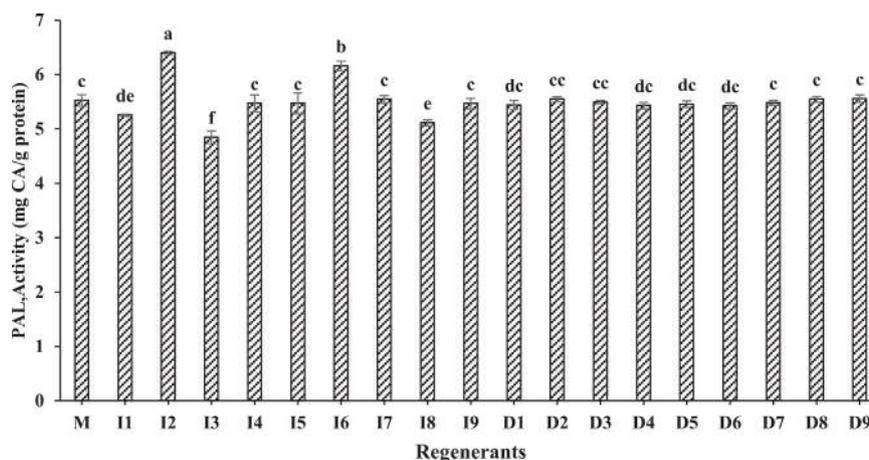
### 3.6. Rosmarinic acid content and PAL activity

Rosmarinic acid (RA) content and phenylalanine ammonia-lyase (PAL) activity of Woodland Sage plantlets were measured at the end of culture using HPLC and spectrophotometric methods, respectively. As shown in Figs. 3 and 4, the activity of the PAL enzyme and the RA content in plantlets acquired from indirect regeneration were different. In contrast, the amount of these parameters in plantlets achieved from direct organogenesis was equal to the mother plant. HPLC analysis of 19 regenerants indicated that I2 and I6 regenerants had the highest RA

**Table 5**

List of ISSR primer sequences with the number and size of amplified fragments generated in direct and indirect organogenesis.

No.	Primers	Primer Sequence	T <sub>m</sub>	Donor Plant		Direct Regeneration			Indirect Regeneration			Range (bp)
				nSB	nTSB	nTPB	P (%)	nSB	nPB	P (%)		
1	UBC-807	(AG) <sub>8</sub> T	50	6	6	0	0	7	2	28.57	250–750	
2	UBC-809	(AG) <sub>8</sub> G	52	4	4	0	0	5	1	20	200–1000	
3	UBC-810	(GA) <sub>8</sub> T	50	6	6	0	0	6	2	33.33	100–800	
4	UBC-811	(GA) <sub>8</sub> C	52	5	5	0	0	5	1	20	250–800	
5	UBC-826	(AC) <sub>8</sub> C	52	5	5	0	0	5	0	0	200–900	
6	UBC-828	(TG) <sub>8</sub> A	50	8	8	0	0	8	0	0	100–900	
7	UBC-834	(AG) <sub>8</sub> YT	52	8	8	0	0	8	1	12.5	100–700	
8	UBC-843	(CT) <sub>8</sub> RA	48	7	7	0	0	7	1	14.28	300–800	
9	UBC-847	(CA) <sub>8</sub> RC	48	4	4	0	0	5	1	20	300–1100	
10	UBC-860	(TG) <sub>8</sub> AA	52	7	7	0	0	8	0	0	100–900	

T<sub>m</sub>: annealing temperature; bp: base pair; R= (A, G), Y= (C, T).**Fig. 2.** - ISSR profiles of the mother plant (M), indirect regenerants (I1-I9) and direct regenerants (D1-D9) of *S. nemorosa* using UBC-847 primer.**Fig. 3.** - PAL activity of the mother plant (M), indirect regenerants (I1-I9) and direct regenerants (D1-D9) of *S. nemorosa*.

content, which was 9.18 and 8.81 mg/g DW, respectively (Fig. 5). Also, I8, I3, and I1 showed the lowest RA content with 6.30, 6.09, and 6.72, respectively. As shown in Fig. 5, cluster analysis grouped the regenerants into three categories, which include group 1 (I2 and I6), group 2 (I1, I3, and I8), and group 3 (I4, I5, I9, and D1–9) (Fig. 6).

#### 4. Discussion

In the present study, direct and indirect organogenesis was induced from *in-vitro* derivative leaves of *S. nemorosa*. It is vital in two respects. Firstly, often true-to-type plants were obtained through direct organogenesis because of direct regeneration from explants (Bhusare et al., 2018). At least the occurrence of genetic changes is crucial in propagating elite genotypes and cultivars (Nayak et al., 2012; Viehmannova

et al., 2016). Secondly, in the indirect regeneration via callus, somaclonal variation may occur. Although it is not desirable for true-to-type plant production, it can be used for genetic improvement and breeding purposes (Bairu et al., 2010; Miguel and Marum, 2011).

In the first experiment of the present research, thirty-five combinations of two cytokinins (BA and TDZ) and two auxins (NAA and 2, 4-D) assessed on organogenesis and callogenesis potential of leaf explants. Explants incubated on PGRs free medium could not stimulate any morphogenesis response. It indicated that the presence of an appropriate plant growth regulator in the *S. nemorosa* culture medium is essential for initiating organogenesis and callus formation. Similar results have been reported in *in-vitro* studies of other *Salvia* species such as *Salvia broussonetii* (Mederos-Molina, 2006), *Salvia canariensis* L. (Mederos-Molina, 2004) and *Salvia miltiorrhiza* (Tsai et al., 2015). Our findings revealed

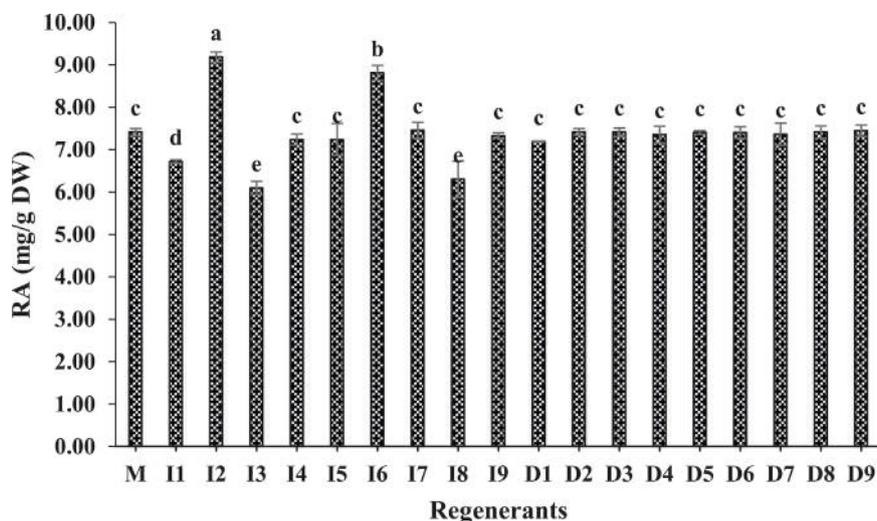


Fig. 4. - Rosmarinic acid content of the mother plant (M), indirect regenerants (I1-I9) and direct regenerants (D1-D9) of *S. nemorosa*.

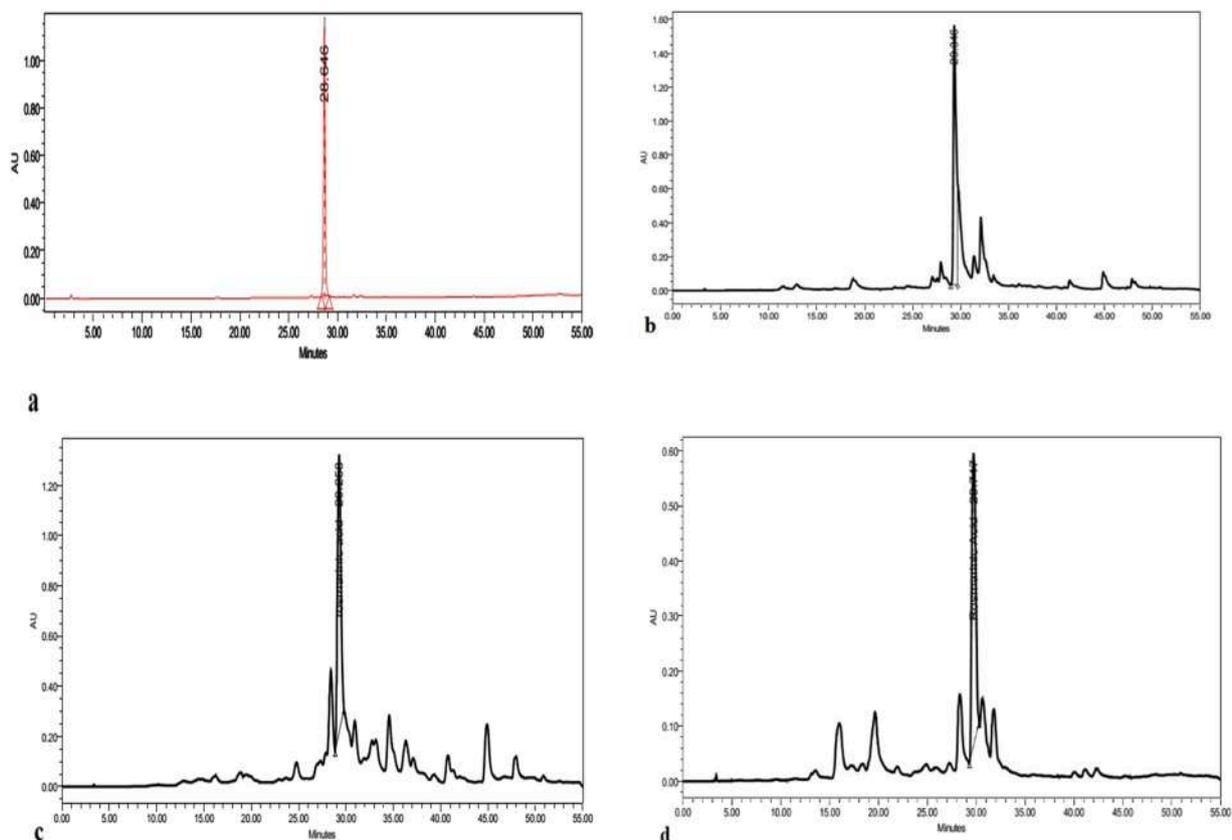


Fig. 5. - HPLC chromatograph of rosmarinic acid standard (a), I2 (b), mother plant (c) and I8 (d) regenerants.

that TDZ solely could provide a satisfactory level of regeneration (37.5–50%), while BA could not induce a shoot or calli without the presence of auxin. Contrary to our results, Skala and Wysokińska (2004) reported that 4.4  $\mu\text{M}$  BA alone provoked direct regeneration from leaf explants of *S. nemorosa*. This difference may be due to diversity in the genotypes studied in the two types of research. In fact, diverse genotypes of one species may exhibit different *in-vitro* responses (Khaliluev et al., 2014; Sabbadini et al., 2019). Thidiazuron is a synthetic plant growth regulator, and it has a potent functionality for morphogenic processes such as shoot organogenesis and proliferation, callus induction, and somatic embryogenesis in tissue culture (Dewir et al., 2018). In the

previous report, the shoot induction of Woodland Sage was achieved by BA as cytokinin and IAA and IBA as auxin (Skala and Wysokińska, 2004).

Our research exhibited that TDZ acted better than BA in high frequency direct and indirect organogenesis in Woodland Sage. Moreover, in this regard, NAA was the better auxin. Some reports suggested that simultaneous use of TDZ and NAA in a culture medium with optimal level enhanced shoot regeneration from leaf explants (Guo et al., 2011). TDZ induces more adventitious shoots than other cytokinins for most of the species, especially in the Lamiaceae family, such as *Salvia x jamensis* (Fraternal et al., 2013), *Salvia miltiorrhiza* (Tsai et al., 2015), *Ajuga bracteosa* (Ali et al., 2018), *Scutellaria bornmuelleri* (Gharari et al., 2019).

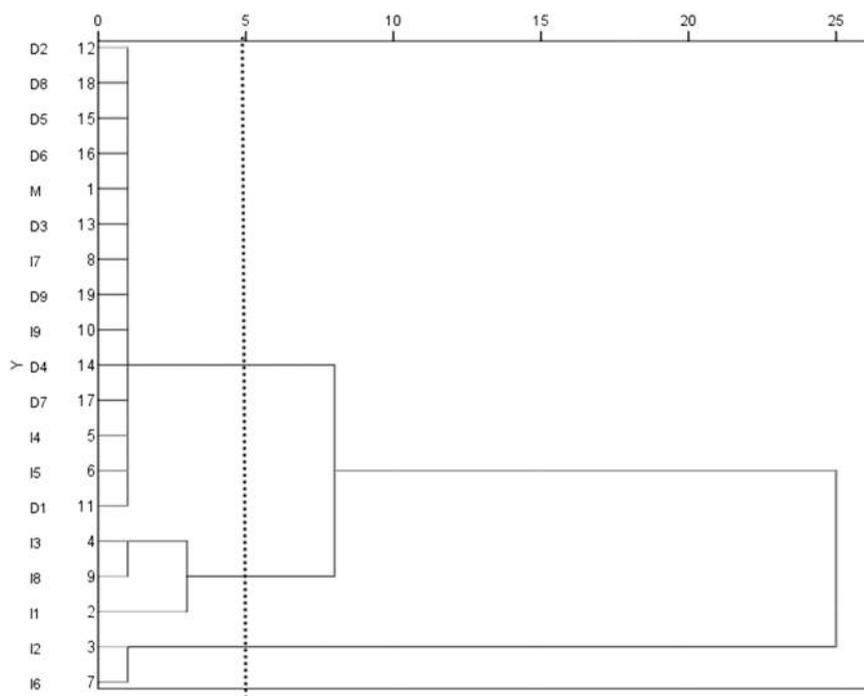


Fig. 6. - Dendrogram of cluster analysis of *S. nemorosa* based on rosmarinic acid content.

TDZ has both auxin and cytokinin activity, and its function relies on concentration and concomitant PGR (Guo et al., 2011). Our findings revealed that TDZ combined with 2, 4-D produced calli and, in combination with NAA, caused shoot regeneration.

According to our findings, 2, 4-D, among all the auxins used in the present experiment, had the best effect on *S. nemorosa* callus formation. As 2, 4-D concentration was raised in the medium, the amount of Woodland Sage callogenesis increased. Also, the presence of a low cytokinin level promoted this performance. In most reports published in callogenesis of *Salvia* genus such as *S. santolinifolia* (Jan et al., 2015, 2018), *S. officinalis* (Lemraski et al., 2014), *S. tomentosa* (Georgiev et al., 2014) and *S. splendens* (Huui et al., 2012), NAA was preferred as auxin source, and it gave better responses. However, Modarres et al. (2018) reported that high levels of 2, 4-D (3 mg/l) along with low levels of Kinetin (1 mg/l) were needed for callus induction in *S. leriifolia*, and it could not respond solely. However, NAA provided a more favorable response as 2, 4-D. These reports contradict our results.

Cytokinins are one of the principal components of the tissue culture medium for shoot multiplication. A proper cytokinin in an optimum concentration appears crucial to this process (Phillips and Garda, 2019). In the present study, attempts were performed to optimize the cytokinin concentration and type to gain healthy multiple shoots with robust growth in *S. nemorosa*. In the earlier report on Woodland Sage published by Skała and Wysokińska (2004), the maximum multiplication rate (5.7 shoots per explant) was achieved in 8.9  $\mu$ M BA, and all levels of BA produce more shoots than TDZ. Despite it, in our study, TDZ, compared to BA, produced the highest shoots at all concentrations. Similar to our results reported on *Salvia x jamensis* (Fraternali et al., 2013). Also, Sharma et al. (2014) evaluated four cytokinin types (BA, TDZ, Kin, and 2iP) on shoot proliferation of *S. splendens* and stated that BA provided the highest multiplied shoots and shoot length compared to others. The consequence of BA than TDZ in shoot proliferation was reported on *S. fruticososa* (Arikat et al., 2004), *S. brachyodon* (Misic et al., 2006), *S. africana-lutea* (Makunga and van Staden, 2007) and *S. guaranitica* (Echeverrigaray et al., 2010) which is in contrast to our results. Overall, we found that adenine sulphate had a better cytokinin activity than the other cytokinins for *S. nemorosa* in vitro multiplication through nodal explants. Compared to TDZ and BA, AdS proliferated fewer shoots, but

they were longer on average. On the other hand, shoots obtained by TDZ and BA exhibited a high level of hyperhydricity (28–54%). While shoots multiplied by AdS did not exhibit hyperhydric symptoms. Adenine sulphate may act synergistically as a cytokinin such as Kinetin or Zeatin, and it is a precursor for enhancing the biosynthesis of natural cytokinins. However, it is possible that AdS also prepares an additional nitrogen source for the cells, which can be taken up more rapidly than inorganic nitrogen. Thus, it can stimulate cell growth, somatic embryogenesis, and callogenesis and considerably enhance the shoot development in tissue culture (Mohd et al., 2014; Naaz et al., 2014). The promotive role of adenine sulphate in shoot proliferation has been described in different species, namely *Pithecellobium dulce* (Goyal et al., 2012), *Cichorium intybus* (Nandagopal and Ranjitha Kumari, 2006), *Althaea officinalis* (Naz and Anis, 2012), *Anacardium occidentale* (Martin, 2003) and *Stevia rebaudiana* (Mohd et al., 2014).

Hyperhydricity is a physiological and morphological abnormality observed in *in-vitro* propagated plants in response to culture conditions, basal medium components, mainly ammonium and plant growth regulator types, and concentrations, especially cytokinins (Liu et al., 2017). In our investigation, the only kind and concentration of cytokinins are diverse in different treatments, and hence, the observed hyperhydric plants were probably due to it. Hyperhydric shoots have higher endogenous cytokinin content than healthy shoots (Ivanova and Staden, 2010). Thidiazuron is more stable in plant tissues than other cytokinins because it is resistant to endogenous cytokinin oxidase. Furthermore, TDZ inhibits cytokinin oxidase activity, which causes an increase in purine cytokinins accumulation in plant tissues (Dewir et al., 2018). Thus, an increase in endogenous cytokinins may explain the high rate of hyperhydricity observed in multiplied shoots of *S. nemorosa* in a medium supplemented with TDZ. In *Pyrus pyrifolia*, Phenylurea derivatives cytokinins like TDZ increased approximately tenfold in the number of hyperhydric shoots than adenine derivatives (Kadota and Niimi, 2003). In several plant species such as *Dianthus caryophyllus* (Genkov et al., 1997), *Aloe polyphylla* (Ivanova and van Staden, 2008), *Cordyline fruticososa* (Dewir et al., 2015), and *Coleonema pulchellum* (Baskaran et al., 2014) TDZ-induced hyperhydricity of regenerated shoots was described.

Shoots incubated on PGRs free medium (control) could not develop adventitious roots at the end of the experiment. Therefore, the presence

of auxin in the rooting medium of the *S. nemorosa* seems necessary. Similar results were reported on other *Salvia* species, namely *S. splendens* (Yu et al., 2015), *S. miltiorrhiza* (Tsai et al., 2015), and *S. canariensis* (Mederos-Molina, 2004). Although, Skala and Wysokińska (2004) reported that *S. nemorosa* shoots could be produced 50% root on PGRs free medium, which does not coincide with our investigation. Our findings revealed that IAA was the better auxin than NAA and IBA in the rooting process because of providing an acceptable root number and length without forming a callus. Calli formation during rooting at the end of the shoot was not desirable because of its hard acclimatization (Sha Valli Khan et al., 1997; Veraplakorn, 2016).

The acclimatization protocol utilized for Woodland Sage *in vitro* raised plantlets was successful, and a 94% survival rate was attained. The present study exhibited a higher survival rate than other *Salvia* species, such as *S. nemorosa* (Skala and Wysokińska, 2004), *S. pratensis* (Ruffoni et al., 2007), *S. sclarea* (Grigoriadou et al., 2020).

To the best of our knowledge, this is the first report on evaluating somaclonal variation in the *in-vitro* culture of *Salvia* species. The ISSR outcomes were obtained from *in-vitro* derived plantlets, and the donor plant showed polymorphic bands. Plants regenerated through direct organogenesis were similar to source plants, and no polymorphic bands were seen. Vice versa, genetic instability has been detected in regenerants obtained by indirect organogenesis. Somaclonal variation is genetic or epigenetic changes found in the cell and tissue cultures, and this term is globally accepted for all forms of tissue culture-derived variants. Several factors, including the source of explants *in-vitro* propagation method, type and concentration of applied plant growth regulators, number and duration of subcultures, and stressful condition of *in-vitro* cultures, may cause somaclonal variation (Bairu et al., 2010; Miguel and Marum, 2011; Wang and Wang, 2012). In the present study, almost all culture conditions were the same, and the observed somaclonal variation was only between the two regeneration methods. Indirect organogenesis is provided by an intermediate callus phase, which is absent in direct organogenesis. Auxins applied during calli induction increased genetic variation by rising DNA-methylation rates (Bairu et al., 2010). In addition, they are often associated with genetic abnormalities such as polyploidy and endoreduplication due to DNA synthesis stimulation (Takagi et al., 2011). Also, 2, 4-D could cause genetic mutations during callus induction (Bouman and De Klerk, 2001). Thus, we can assume that maybe genetic changes during callus initiation are the main reason for the somaclonal variation observed in the indirect regeneration of Woodland Sage. For several plant species, such as *Titanotrichum oldhamii* (Takagi et al., 2011), strawberry (Biswas et al., 2009), *Chirita swinglei* (Chen et al., 2016), *Aloe vera* (Rathore et al., 2011) and *Bletilia striata* (Wang and Tian, 2014) somaclonal variation was observed in indirect regeneration. In comparison, direct regenerants showed genetic fidelity. Advantage of ISSR markers in the examination of genetic stability and somaclonal variation of *in-vitro* derived plantlets, similar to works on crops such as *Vanilla planifolia* (Pastelin Solano et al., 2019), *Puya berteroniana* (Viehmanna et al., 2016), *Withania coagulans* (Rathore et al., 2016), *Artemisia absinthium* (Kour et al., 2014) and *Moringa peregrina* (Al Khateeb et al., 2013) were validated by our findings.

In the current study, there was a significant difference in the activity of the PAL enzyme among indirect regenerants. In contrast, PAL activity was almost the same in direct regenerants. PAL is a branch point enzyme between the primary and the secondary metabolism, and is a key enzyme for regulating the influx of phenylalanine to the biosynthesis of phenolic compounds (Dong et al., 2010). Evidence has revealed that genetic, epigenetic changes, and external factors such as stress and elicitors can be affected PAL enzyme activity (Heydari et al., 2020b). In addition, several studies have confirmed that rosmarinic acid production directly correlates with PAL enzyme activity (Dong et al., 2010; Hou et al., 2020; Li et al., 2020). As a result, it can be concluded that the stability of PAL enzyme activity and the accumulation of rosmarinic acid among directly regenerated plants in the present study might explain the lack of genetic and epigenetic changes during direct organogenesis from

*S. nemorosa* leaf explants. Also, according to the ISSR analysis results, the difference in the amount of rosmarinic acid production in indirect regenerants can be explained by genetic changes. Rosmarinic acid is a phenolic molecule with various applications, varying from food preservatives to cosmetics, and is also a molecule of interest to the pharmaceutical industry (Marchev et al., 2021). The RA possesses anti-inflammatory, anti-oxidative, antibacterial, antiviral, anti-hyperglycemic, anticancer, hepatoprotective, analgesic, cardioprotective, and neuroprotective properties. It is used in modern medicine to treat Alzheimer's disease, diabetes, fibrosis, and various types of cancer such as colorectal, breast, pancreatic, and lung cancers (Hitl et al., 2021; Trócsányi et al., 2020). Rosmarinic acid (RA) is the most abundant phenolic compound detected in Woodland Sage (Bahadori et al., 2017). Salvianolic acids, which have been used to treat cardiovascular disorders, fibrosis, and cancer, are another significant phenolic compound isolated from *S. nemorosa* (Heydari et al., 2020b).

## 5. Conclusion

The current study described an efficient and high-frequency *in vitro* regeneration protocol from leaf explants of *S. nemorosa* through direct and indirect organogenesis. The investigations revealed that the types, combinations, and ratios of plant growth regulators (PGRs) significantly affected direct organogenesis and callogenesis responses. TDZ, in combination with NAA, induced maximum direct adventitious shoot organogenesis from leaf explants, and by increasing the NAA ratio, the organogenesis rate decreased. The highest callus induction was observed when 2, 4-D was combined with BA or TDZ at a 4:1 ratio, and callogenesis decreased as this ratio declined. Our finding showed that TDZ produced maximum adventitious shoot during the proliferation stage, but most of them (40–50%) showed hyperhydric symptoms. Thus, adenine Sulphate is preferred as a PGR with cytokinin activity because it proliferated adventitious shoots without hyperhydricity. The plantlets rooted well in all the tested auxin. All the rooted plantlets were acclimatized with 94% survival in field conditions. For the first time, molecular and biochemical markers have been employed in present study to investigate somaclonal variations in *Salvia* species *in vitro* regeneration. ISSR and biochemical analysis confirmed the true-to-type nature of direct regenerated plantlets. Moreover, these analyses confirmed that somaclonal variation occurred during indirect organogenesis through the callus. The present direct organogenesis protocol is highly efficient and suited for multiplication and transgenic studies to improve the ornamental and medicinal traits of Woodland Sage. Furthermore, the indirect organogenesis protocol may promote the selection of somaclonal variants, which may help select ornamentally/ pharmaceutically important variants of commercial value.

## CRedit authorship contribution statement

**Hamid Reza Heydari:** Data curation, Methodology, Software, Investigation, Writing - original draft, Writing - review & editing.  
**Esmail Chamani:** Conceptualization, Methodology, Supervision, Writing - review & editing.

## Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

## Data availability

Data will be made available on request.

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