Gold nanoparticles and electromagnetic irradiation in tissue culture systems of bleeding heart: biochemical, physiological, and (cyto) genetic effects

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Abstract
This study aimed to analyze the effect of various mutagens on the in vitro development, physiological activity, acclimatization efficiency, and genetic integrity of Lamprocapnos spectabilis ‘Valentine’. Gold nanoparticles (AuNPs), microwaves, and X-rays were used at different doses. The profiles of primary and secondary metabolites and the enzymatic activity in the produced plants were studied. The usefulness of various genetic markers in the detection of mutations in the species was compared. The genome size of L. spectabilis was estimated for the first time. It was found that the addition of AuNPs into the culture medium had a positive impact on the in vitro development and multiplication of plants. All of the shoots regenerated adventitious roots, but plants subjected to the longest microwave irradiation (3 × 9 s) and the non-treated control had the lowest acclimatization efficiency. Application of mutagens significantly affected the activity and profile of most enzymes and phytochemicals studied, however, the final effect depended on the agent type and dose. Mutations were detected by DAMD, RAPD, and SCoT markers in 7.5% of plants, but not by ISSRs. Phenotype variation in leaf shape was found in four plants. The genome size of L. spectabilis was found to be very small; about 1281 Mbp.

Key message
Gold nanoparticles improve the micropropagation and acclimatization efficiencies in bleeding heart—a species of very small genome size. X-rays, on the other hand, are the most suitable for inducing phenotypical changes. Microwaves are the least useful for both propagation and breeding purposes.

Keywords Genetic variation · Genome size · Hormesis · Lamprocapnos spectabilis (L.) Fukuhara · Micropropagation · Mutation breeding · Spectral fingerprinting

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Introduction

*Lamprocapnos spectabilis* (L.) Fukuhasa is a rhizomatous herbaceous perennial, native to Siberia, northern China, Korea, and Japan (Kamińska et al. 2004; Cho 2018). It is also known as bleeding heart, fallopian buds, lyre flower, or lady-in-a-bath. The species is placed in the Fumarioidae, a subfamily of the poppy family—Papaveraceae (Park et al. 2018). It is the only species in the monotypic genus *Lamprocapnos* but is still widely referenced under its former name *Dicentra spectabilis* (L.) Lem. (syn. *Fumaria spectabilis* L.). Bleeding heart is appreciated in landscape architecture, gardens, and floristry for its decorative leaves and long-viable heart-shaped red, pink, and white flowers, produced in spring. It is also valued in the pharmaceutical and cosmetic markets due to the presence of biologically active isoquinoline alkaloids (Adamski et al. 2020), such as protopine and sanguinarine (Och et al. 2017). Extracts of bleeding heart showed cytotoxic activity against the tested human squamous carcinoma and adenocarcinoma cells (Petruczynik et al. 2019). Therefore, the species is recently focusing the attention of biotechnologists, although its biology and characteristic (e.g. stress-related physiological reactions) are not fully understood and require more research.

Conventionally, *L. spectabilis* is propagated vegetatively through softwood cuttings and root division (mainly) or, less frequently, by seeds (Hodges 2012). The viability of seeds is low and they must be sown while fresh, immediately after collection (Deno 1993). Next, the seeds must undergo dormancy through winter to germinate in the spring of the following year. Therefore, there is a long period (almost 1 year) between sowing and germination (Cho et al. 2020). Moreover, the generative method of reproduction fails to retain the characteristics of the mother plant and fulfill the demand of the pharmaceutical market. Cuttings and division provide plants true to the parent plant and a quicker bloom time, however, their yield is limited and climate-dependent (Lema-Rumińska et al. 2019). Hence, it is essential to make novel efforts to establish sufficient plants of *L. spectabilis* by artificial propagation and breeding methods to meet the elevated market demand.

In vitro tissue culture systems have been successfully utilized for propagation, breeding, preservation, and secondary metabolite production in numerous ornamental and medicinal plant species in the past decades (Rout and Jain 2020). However, in some species, classical plant growth regulators (PGRs), such as benzyl adenine (BA) cause certain morpho-physiological, anatomical, and biochemical disorders that negatively affect the quality of the produced biological material, particularly during acclimatization (Bidabadi and Jain 2020; Manokari et al. 2021). Bleeding heart can be classified as a species recalcitrant to in vitro propagation as the simultaneous exogenous supply of common auxins and cytokinins into the culture medium, even at low concentrations, results in the abundant secretion of harmful polyphenols, formation of callus, and somaclonal variation occurrence (Kulus et al. 2020). Moreover, the species does not produce efficiently adventitious shoots from leaf, petiole, or internode explants (Kulus and Tymoszuk 2020). Therefore, other factors stimulating the in vitro development of this species should be recognized.

Breeding programs lead to an increase in the number of available cultivars of bleeding heart in the twentieth and twenty-first centuries (Hodges 2012). In vitro techniques were established with the intent to develop commercial cultivars of this species via genetic transformation and regeneration from somatic embryos (Lee and Lee 2003). The site-directed CRISPR/Cas-based technologies also gain increasing interest in the breeding of essential crop plants (Miler et al. 2021). Genetic engineering, however, is a complex, time-absorbing, and expensive method of plant improvement. It is also the subject of public debate. Classical mutation breeding, on the other hand, based on the concept of random mutation induction, is an attractive solution to these problems and it remains still an important method for breeding ornamental plants alongside hybridization (Holme et al. 2019; Datta 2020). The method allows broadening the scope of genetic variation without profoundly changing the whole genome of valued individuals and with no need for sophisticated molecular laboratories equipped with expensive devices, and employees trained in genetic engineering (Jo and Kim 2019). Moreover, it is universal and does not require deep knowledge of genes sequences, structures, and functions (Shelake et al. 2019).

Mutagenic factors used in plant breeding can be classified into two groups: chemical and physical. Among the first group, ethyl methanesulphonate is deployed most often, however, recent studies have shown that nanoparticles (NPs) can be efficacious to induce genotoxic effects in plants, due to their ease of interacting with plant cells (Tymoszuk and Miler 2019; Miler et al. 2021). Silver nanoparticles (AgNPs) are the best-studied in terms of genetic variation induction and catalysis of reduction–oxidation reactions in a plant cell (Tymoszuk and Kulus 2020). As opposed to AgNPs, fewer studies determining the influence of gold nanoparticles (AuNPs) on plants have been published (Kulus and Tymoszuk 2021). Physical mutagens, such as X- or gamma-rays, and heavy-ion beams are more eco-friendly as their deployment does not result in chemically harmful waste production. Unfortunately, the accessibility of the devices used for irradiation treatment can be a limiting factor for plant breeders, as those appliances are usually owned by national atomic agencies or medical/scientific institutes (Miler et al. 2019).
Alternatively, a common microwave oven can be an inexpensive and effective source of mutagenic radiation (microwaves; MW), although, to date, this approach has bee used exclusively with *Chrysanthemum × morifolium* (Ramat.) Hemsly. by Miler and Kulus (2018).

Recognizing the extent of genetic variation is a key factor for any effective genetic improvement program, micropropagation protocol, and also for any successful hybridization plan to obtain progenies with desirable traits (Adhikari et al. 2020). Due to their simplicity, rapidity, and versatility, genotyping methods based on PCR are now commonly utilized to assess the genetic variation in agricultural, medicinal, and horticultural plants. Single primer amplification reaction (SPAR) methods; such as Randomly Amplified Polymorphic DNA (RAPD) (Williams et al. 1990), Directed Amplification of Minisatellite DNA (DAMD) (Heath et al. 1993), Inter Sequence Simple Repeat (ISSR) (Zietkiewicz et al. 1994), and Start Codon Targeted Polymorphism (SCoT) (Collard and Mackill 2009); are considered powerful tools for studying genetic diversity in plants, especially in those species that have not been well-studied or those for which fingerprinting keys had not been developed. Although each SPAR method has its drawbacks, cumulative data support a comprehensive elucidation of the genetic variability within a species (Nadeem et al. 2017; Adhikari et al. 2020).

Flow cytometry (FCM), chemotaxonomy, and phenotyping can be additionally used to obtain a complete view of the stability of plant material. Flow cytometric evaluation of nuclear DNA content is a relatively simple, fast, and low-cost method of verifying genome size stability, requiring only a small amount of biological material (Sliwinska 2018). Changes in metabolic profiles can be studied with the help of chemical fingerprinting. Chemotaxonomy; i.e. the analysis of confirmable differences in the biochemical composition of roots, leaves, flowers, and fruit; is a supplementary method of cultivar identification. Variation in the color of plant organs gave a basis for demonstrating differences across cultivars of various plant species (Lema-Rumińska et al. 2018). Moreover, chemical portrait given by spectral fingerprinting provides information valuable to the pharmaceutical industry.

The literature survey reveals no information of mutation breeding or genome size estimation in bleeding heart. Therefore, the present study was performed (i) to study the impact of chemical (AuNPs) and physical agents (MW and X-rays), applied at various doses, on the in vitro development, micropropagation efficiency, and post-acclimatization quality of bleeding heart ‘Valentine’; (ii) to analyze the enzymatic activity and phytochemicals content in the shoots produced; (iii) to assess for the first time the genome size of *Lamprocapnos spectabilis* using FCM; (iv) to analyze the genetic stability of the greenhouse-grown plants using PCR-based markers; (v) to determine the usefulness of various SPAR marker systems in the detection of genetic variation in bleeding heart.

### Materials and methods

#### Plant material and in vitro culture conditions

Single-node shoot fragments of *Lamprocapnos spectabilis* (L.) Fukuhara ‘Valentine’, isolated from 8-week-old microcuttings, were vertically inoculated in 350-mL glass jars filled with 40 mL of MS (Murashige and Skoog 1962) medium augmented with 1.0 mg L\(^{-1}\) kinetin (Sigma-Aldrich, St. Louis, MO, USA). The medium was additionally supplemented with 30 g L\(^{-1}\) of sucrose and solidified with 8.0 g L\(^{-1}\) agar (Biocorp, Warsaw, Poland). After adding all the components, the pH was adjusted to 5.8 before the medium was autoclaved at 121 °C and 105 kPa for 20 min. Nine explants were placed in each culture jar sealed with a plastic cap. One jar was considered a single repetition.

The cultures were grown at 24 ± 1 °C and exposed to a 16-h photoperiod. Standard cool daylight was provided by TLD54/36 W fluorescent lamps (Koninklijke Philips Electronics N.V., Eindhoven, the Netherlands) with the photosynthetic photon flux density of approximately 35 \(\mu\)mol m\(^{-2}\) s\(^{-1}\) (Fig. 1A).

#### Treatment with chemical and physical agents

Immediately after inoculation in the culture medium, the explants were treated with chemical (AuNPs) or physical agents (MW and X-rays).

#### AuNPs treatment

AuNPs, spherical in shape, of 13 ± 3 nm diameter and at 50, 75, or 100 ppm concentration, obtained from Nanoparticles Innovation NPIN s.c. (Łódź, Poland), were cold-sterilized through a syringe filter (Minisart® RC 25, pore size 0.20 μm; Sartorius AG, Göttingen, Germany) and poured onto the culture medium, 2 mL per one culture jar (Fig. 1B). The detailed characteristics and synthesis protocol of nanoparticles are given in Kulus and Tymoszek (2021). AuNPs were synthesized using a chemical reduction method. Briefly, an aqueous solution of gold (III) chloride hydrate was heated to the boiling point under reflux and, then, an aqueous solution of sodium citrate was added to the reaction mixture. The solution was heated for 15 min under stirring, and next the colloid was cooled down to room temperature.
Microwave treatment

SAMSUNG MW-73B-S microwave oven with the power of 800 W cm\(^{-2}\) and the frequency of 2.45 GHz was used as the source of MW radiation (Fig. 1C). The explants were irradiated with MW for 9 s in repeated cycles, as following: 1×9 s, 2×9 s (a total of 18 s), and 3×9 s (a total of 27 s). To maintain a constant room temperature between the successive irradiations, the jars with the explants were cooled in a water bath at the temperature of 10 °C for 1 min.
Irradiation with X-rays

For the time of X-ray treatment, jars with inoculated nodal segments were filled with sterile distilled water so the explants were immersed about 5 mm below the water surface to avoid the usual confusion arising from a lack of exact dose determination. The objects were irradiated with X-rays of 6MV Nominal Accelerating Potential (NAP) at the Oncology Center in Bydgoszcz, Poland. A conventional medical accelerator Clinac 2300CD Silhouette from Varian Medical Systems (Palo Alto, Santa Clara, CA, USA) was used for the experiment (Fig. 1D). Three total doses were delivered to the explants, i.e. 10 Gy or 15 Gy or 20 Gy, with the dose rate 3.39 Gy min−1 (300 MU min−1). The source-surface distance (SSD) was 90.8 cm.

Nodes not treated with AuNPs, MW, or X-rays were considered as the control. Each treatment was performed in five repetitions, i.e. with 45 explants in every experimental object. A total of 450 nodal segments were used.

Biometric analysis of in vitro-grown plants

After 10 weeks of in vitro culture, the share of viable explants, the number of shoots produced, propagation ratio, and spontaneous rooting ability were determined. The propagation ratio was assessed by counting the number of secondary explants (nodal segments) that could be excised from the recovered shoots. Foliar biomass; i.e. the fresh (FW), turgid (TW), and dry (DW) weight of leaves were determined to calculate the relative water content (RTW) in plants based on Dastborhan and Ghassemi-Golezani (2015) with minor modifications. Three young leaves from six plants from each experimental object were used in the analysis with one plant being a single repetition.

Spectral fingerprinting

The spectral analysis of enzymes and pigments was performed 10 and 11 weeks after culture initiation using 200- and 100-mg fresh tissue samples, respectively, in six repetitions and a spectrophotometer SmartSpec PlusTM (BioRad, Hercules, CA, USA).

Determination of enzymatic activity

Samples taken from in vitro-grown shoots were homogenized in 100 mM phosphate buffer (pH 7.4) containing 1 mM EDTA, 1 mM dithiothreitol (DTT), and 2% polyvinylpyrrolidone (PVP) (all chemicals provided by Chemat, Gdańsk, Poland), according to Homae and Ehsanpour (2016). The homogenates were centrifuged at 13,000×g for 20 min at 4 °C (Centrifuge MPW-260R, MPW MED INSTRUMENTS, Warsaw, Poland) and supernatants were used to determine the activities of antioxidant enzymes and protein content.

Glutathione reductase (GR; E.C. 1.8.1.7) activity was determined spectrophotometrically by measuring the enzymatic-dependent oxidation of NADPH (Mallebrera et al. 2014). One unit of GR was defined as the amount of enzyme that oxidized 1.0 μmol of NADPH in one minute. The guaiacol peroxidase (G-POD; E.C. 1.11.1.7) and ascorbate peroxidase (APX; E.C. 1.11.1.11) activities were monitored based on the polymerization of guaiacol into tetraguaiacol and ascorbate consumption according to Maehly and Chance (1954) and Nakano and Asada (1981), respectively, with modifications described by Nowogór ska and Patykowski (2015). Protein content was measured based on the Bradford method (1976) with bovine serum albumin (BSA) as the standard. The spectrophotometric analysis of extracts was performed at specific wavelengths: for proteins at 595 nm, for GR at 340 nm, for G-POD at 470 nm, and for APX at 290 nm. The enzymatic activity U [μmol min−1] was calculated per 1 mg of protein. Next, the enzyme unit U was converted to katal, with the assumption that 1 U = 1/60 μkat = 16.67 nkat.

Analysis of pigments content

Chlorophylls and carotenoids (carotenes and xanthophylls) were extracted with the procedure described by Bulda et al. (2008) using a 4:1 mixture of petroleum ether (PE, fraction 60–90) and tetrahydrofuran (THF) and 100 mg of fresh tissue samples. The total flavonoid content of methanol extract was determined by the aluminum chloride method (Brighente et al. 2008). The flavonol contents content was evaluated with the method described by Kabir et al. (2016). Anthocyanins were extracted using 200 mg leaf samples and methanol containing 1% HCl, according to the Harborne (1967) method. The same extract was used to analyze the total phenolic and tannin concentrations with the Folin–Ciocalteau reagent, according to the Waterhouse (2001) and Kabir et al. (2016) procedures, respectively. The total flavonoid, flavonol; phenolic; and tannin contents were calculated using quercetin; gallic acid; and tannic acid as the calibration standards (0–400 mg L−1), respectively.

The spectrophotometric analysis of extracts was performed at specific wavelengths (λ max): for flavonoids at 420 nm, for flavonols at 440 nm (maximum absorbance of 3-hydroxy flavone), for carotenoids at 480 and 495 nm (β-carotene and lutein), for anthocyanins (cyanidin-3-glucoside) at 530 nm, for chlorophyll a and b at 645 and 655 nm, and for tannins and phenolics at 725 and 765 nm, respectively. The content of metabolites was calculated per gram of fresh matter.

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Estimation of nuclear DNA content in in vitro-derived plantlets by flow cytometry

The fresh young leaves of each 12-week-old shoot were sampled to measure the 2C DNA content. The samples for the FCM analysis were prepared according to the procedure described by Rewers et al. (2012), using 1 mL of nuclei-isolation buffer (200 mM Tris; 4 mM MgCl₂ × 6H₂O; 0.5% Triton X-100; pH 7.0; Pfosser et al. 1995), supplemented with propidium iodide (PI; 50 μg mL⁻¹) and ribonuclease A (RNase A; 50 μg mL⁻¹). *Solanum lycopersicum* L. ‘Stupicke’ (2C = 1.96 pg; Dolezel et al. 1992) served as an internal standard. For each sample, at least 7000 nuclei were analyzed using a CyFlow SL Green (Partec GmbH, Münster, Germany) flow cytometer, equipped with a high-grade solid-state laser with green light emission at 532 nm, long-pass filter RG 590 E, DM 560 A, as well as with side (SSC) and forward (FSC) scatters. Analyses were performed on 15 biological replicates using linear amplification. Histograms were evaluated using FloMax software (Partec GmbH, Münster, Germany). The coefficient of variation (CV) of the G₀/G₁ peak of *L. spectabilis* ranged between 2.56 and 4.09%. Nuclear DNA content was calculated using the linear relationship between the ratio of the 2C peak positions of bleeding heart/internal standard on the histogram of fluorescence intensities. The 2C nuclear DNA content in picograms (pg) was transformed into megabase pairs (Mbp) of nucleotides, using the conversion 1 pg = 978 Mbp (Doležel et al. 2003).

In vitro rooting, acclimatization, and ex vitro growth

Twenty-five to fifty shoots from each experimental combination were rooted and acclimatized. For this purpose, the shoots 3 cm in length were dissected and transferred on the rooting MS medium containing 2.0 mg L⁻¹ of indole-3-acetic acid (IAA) for 21 days.

An additional group of 15 shootlets (three per repetition) representing every experimental combination was grown on the same medium for 6 weeks to collect data on the architecture of their root systems. Root systems were excised from the shoots and scanned (Epson STD4800 scanner, Suwa, Japan) in the transparent polypropylene cuvettes filled with water. Next, root system pictures were analyzed and a mean total length of the roots, a mean root system area, and root diameter, as well as the number of forks and tips, were measured using the imaging software WinRHIZO™ (Reagen Instruments, Quebec, Canada).

Acclimatization was conducted from October 26 to November 13 in natural light conditions in a glasshouse in five replications (with 5–10 plants per repetition). Bleeding hearts were grown individually in plastic trays filled with a mixture of peat and perlite (2:1), sprayed with water, and covered with perforated foil and geo-cover. The acclimatization effectiveness, i.e. survival rate and quality of plants, was assessed after 42 days. The survival was calculated as the percentage of the number of living plants out of the initial number of acclimatized bleeding hearts. Plants that remained in the healthy form were considered fair quality. Those, which were tiny or had withered leaves were considered poor quality. The quality of shoots with evident signs of growth, i.e. development of new healthy leaves, was considered superb.

Analysis of genetic fidelity

The genetic fidelity of plants was assessed using DAMD, ISSR, RAPD, and SCoT marker systems, after 6 weeks of ex vitro growth. A total of 80 shoots (eight from each experimental object) were analyzed.

Total genomic DNA was isolated from fresh tissues using a FastPrep-24 5G bead beating grinder and lysis system (MP Biomedicals Irvine, CA, USA) and Genomic Mini AX Plant Spin kit (A&A Biotechnology, Gdynia, Poland), according to the manufacturer’s instruction.

Five primers from each marker system were used in the PCR reaction. PCR was performed in a gradient C1000 Touch thermal cycler with heated cover (Bio-Rad, Hercules, CA, USA) in the 25-μL reaction solution. The composition of the reaction solution, PCR profiles, and electrophoretic separation of amplified DNA fragments were described in Adhikari et al. (2020 with modifications) and Kulus (2020). Gel images were recorded using a GelDoc XR + Gel Photodocumentation System (Bio-Rad, Hercules, CA, USA) UV transilluminator with Image Lab 4.1 software. Molecular weights of the fragments were estimated using a 100–5000 bp DNA molecular marker (Gene Ruler TM Express DNA Ladder, Thermo Fisher Scientific, Waltham, MA, USA).

The banding patterns were recorded as 0–1 binary matrix, where “1” indicates the presence and “0” the absence of a given fragment followed by statistical analysis. For every primer tested, the numbers of monomorphic, polymorphic, and specific (unique) bands were counted. Values of Heterozygosity index (H), Polymorphic Information Content (PIC), Effective multiplex ratio (E), Marker Index (MI), Discriminating power (D), and Resolving power (R) were also investigated for every primer and marker system used (Amiryousefi et al. 2018).

Statistical analysis

The experiment was set up in a completely randomized design. The experimental data were presented as mean ± standard error (SE). To obtain the normal distribution of the values expressed as a percentage, the Freeman–Tukey double-arcsine transformation was used. The
results were evaluated by the one-factor analysis of variance (ANOVA) and the Duncan’s Multiple Range Test at \( p = 0.05 \) using Statistica 12.0 (Statsoft, Cracow, Poland). Tables with results provide real, untransformed numerical data, with the alphabet indicating the homogeneous groups. GenAlex 6.5 software package (Peakall and Smouse 2012) was used to count the genetic distance, perform the principal component analysis (PCoA) and run the analysis of molecular variance (AMOVA) with the assumption that the experimental objects (three factors used at three levels, each and the non-treated control) are 10 separate populations. iMEC software was used to calculate the polymorphism indices (Amiryousefi et al. 2018).

**Results**

**In vitro multiplication and biometric analysis of bleeding heart**

Neither gold nanoparticles nor electromagnetic irradiation affected the viability of the explants, which reached 86–100% (Table 1). Explants treated with AuNPs produced significantly more shoots (2.2–4.2) than the untreated control (1.1). The propagation ratio was also positively affected by nanoparticles, with 22.9 secondary explants produced after adding 75 ppm AuNPs into the culture medium (Table 1; Fig. 1E). On the other hand, X-irradiation had a deleterious effect on spontaneous rooting (24.0–41.4% of shoots) compared to the control (80.0%). No evident callus formation was reported in any experimental object.

| Treatment | FW (mg) | DW (mg) | RWC (%) |
|-----------|---------|---------|---------|
| Control   | 5.19±0.55 bc* | 1.27±0.22 b-d | 88.68±2.06 a |
| Gold nanoparticles | 50 ppm | 4.75±0.37 bc | 1.01±0.10 d | 90.68±3.27 a |
| 75 ppm | 3.86±0.22 c | 1.39±0.10 b-d | 88.88±2.05 a |
| 100 ppm | 4.58±0.50 bc | 1.01±0.07 d | 85.46±2.67 a |
| Microwaves | 9 s | 5.22±0.29 bc | 1.17±0.05 cd | 84.42±1.96 a |
| 18 s | 4.69±0.41 bc | 0.98±0.10 d | 95.58±1.27 a |
| 27 s | 3.68±0.53 c | 0.96±0.16 d | 90.46±2.51 a |
| X-rays | 10 Gy | 6.46±0.90 ab | 1.59±0.22 bc | 91.82±2.31 a |
| 15 Gy | 7.49±0.85 ab | 2.27±0.27 a | 87.87±1.89 a |
| 20 Gy | 7.66±0.81 a | 1.79±0.22 ab | 93.03±1.57 a |

*Values are means ± SE, different letters show significant differences using Duncan’s test (\( P \) value < 0.05)

Irradiation with X-rays, especially 15 Gy and 20 Gy, increased the fresh and dry weights of leaves compared to the control (Table 2). In contrast, there was no influence of the factors studied on the relative water content in the plants (85.46–95.58%).

**Table 1** Influence of chemical and physical agents on the survival and morphogenic response in *L. spectabilis* ‘Valentine’ after 10 weeks of in vitro culture

| Treatment         | Survival (%) | No. of shoots | Propagation ratio | Rooting (%) |
|-------------------|--------------|---------------|--------------------|-------------|
| Control           | 94.3 a*      | 1.1±0.1 d     | 7.9±0.5 c          | 80.0 ab     |
| Gold nanoparticles|              |               |                    |             |
| 50 ppm            | 100 a        | 2.2±0.2 bc    | 19.8±0.7 a         | 57.1 bc     |
| 75 ppm            | 100 a        | 4.2±0.5 a     | 22.9±2.5 a         | 74.3 ab     |
| 100 ppm           | 95.6 a       | 2.7±0.3 b     | 18.6±1.9 b         | 55.8 bc     |
| Microwaves        |              |               |                    |             |
| 9 s               | 100 a        | 1.9±0.7 b-d   | 8.1±0.6 c          | 78.0 ab     |
| 18 s              | 100 a        | 2.0±0.1 b-d   | 11.2±0.6 c         | 83.3 a      |
| 27 s              | 100 a        | 1.8±0.3 b-d   | 10.7±1.1 c         | 80.5 ab     |
| X-rays            |              |               |                    |             |
| 10 Gy             | 94.5 a       | 1.5±0.1 cd    | 9.8±0.5 c          | 38.0 cd     |
| 15 Gy             | 86.0 a       | 1.7±0.2 b-d   | 8.9±0.4 c          | 41.4 cd     |
| 20 Gy             | 100 a        | 1.6±0.1 b-d   | 9.0±0.5 c          | 24.0 d      |

*Values are means ± standard errors (SE), different letters show significant differences using Duncan’s test (\( P \) value < 0.05)
Metabolic and physiological activity of bleeding heart

Plants exposed to 18–27 s microwave treatment and 20 Gy of X-rays had an over two-fold higher APX activity (2.08–2.68 nkat) than the control (0.93 nkat; Table 3). Most of the experimental objects showed higher G-POD activity (2.06–3.54 nkat) compared to the untreated control (0.75 nkat), except for those exposed to 50 and 75 ppm AuNPs, and 9 s microwaving. As for GR, only shoots produced on the medium with the highest AuNPs concentration differed significantly from the control (328.88 and 159.14 nkat, respectively).

The highest content of chlorophyll $a$ (268.7 µg g$^{-1}$ FW), chlorophyll $b$ (284.9 µg g$^{-1}$ FW), and total chlorophyll (553.6 µg g$^{-1}$ FW) was found in the shoots produced after 18 s of MW treatment (Table 4). Microshoots produced in the medium with 100 ppm AuNPs, on the other hand, contained less total chlorophyll (217.9 µg g$^{-1}$ FW) than the untreated control (295.4 µg g$^{-1}$ FW). Nonetheless, none of the experimental objects differed in terms of chlorophyll $a/b$ ratio from the control object. The highest content of carotenoids, xanthophylls, and total carotenoids was also found

### Table 4 Influence of chemical and physical agents on the content of chlorophyll $a$ (Cl $a$), chlorophyll $b$ (Cl $b$), total chlorophyll (Ct), carotenes, xanthophylls, and carotenoids in *L. spectabilis* ‘Valentine’ after 11 weeks of in vitro culture

| Treatment   | Cl $a$ (µg g$^{-1}$) | Cl $b$ (µg g$^{-1}$) | Ct (µg g$^{-1}$) | Cl $a/b$ | Carotenes (µg g$^{-1}$) | Xanthophylls (µg g$^{-1}$) | Carotenoids (µg g$^{-1}$) |
|-------------|-----------------------|-----------------------|------------------|----------|-------------------------|---------------------------|---------------------------|
| Control     | 167.5 ± 11.9 c-e*     | 127.9 ± 10.7 b-d      | 295.4 ± 8.0 bc   | 1.39 ± 0.2 a-d | 43.8 ± 7.8 a          | 66.5 ± 10.6 bc            | 110.3 ± 7.6 b             |
| Gold nanoparticles  
50 ppm  | 148.5 ± 9.6 de         | 170.0 ± 13.3 b        | 318.5 ± 22.5 bc  | 0.88 ± 0.0 d  | 31.5 ± 5.4 ab          | 85.9 ± 5.2 bc             | 117.4 ± 7.6 b             |
| 75 ppm  | 142.7 ± 13.9 de        | 143.1 ± 11.9 bc       | 285.9 ± 9.9 cd   | 1.07 ± 0.2 b-d | 8.3 ± 1.2 c           | 84.8 ± 7.6 bc             | 93.0 ± 9.5 b              |
| 100 ppm | 123.0 ± 9.3 e          | 94.8 ± 10.8 d         | 217.9 ± 7.1 d    | 1.30 ± 0.3 a-d | 27.4 ± 7.3 a-c         | 62.3 ± 8.8 e              | 89.7 ± 8.0 b              |
| Microwaves  
9 s       | 219.2 ± 45.1 a-c       | 155.9 ± 26.6 bc       | 375.0 ± 60.5 b   | 1.46 ± 0.3 a-d | 23.8 ± 8.8 a-c         | 89.3 ± 11.8 bc            | 113.1 ± 16.5 b            |
| 18 s       | 268.7 ± 12.3 a         | 284.9 ± 21.6 a        | 553.6 ± 15.6 a   | 0.99 ± 0.1 cd | 41.2 ± 6.8 a          | 175.6 ± 11.1 a            | 216.8 ± 15.5 a            |
| 27 s       | 156.9 ± 14.4 c-e       | 154.5 ± 6.5 bc        | 311.4 ± 19.2 e   | 1.01 ± 0.1 b-d | 12.1 ± 3.8 bc         | 91.6 ± 4.5 b              | 103.7 ± 3.7 b             |
| X-rays     
10 Gy     | 203.8 ± 4.8 b-d        | 134.6 ± 11.8 b-d      | 338.4 ± 8.5 bc   | 1.58 ± 0.2 a-c | 26.5 ± 3.8 a-c         | 75.7 ± 9.4 bc             | 102.2 ± 10.7 b            |
| 15 Gy     | 184.0 ± 26.4 b-e       | 114.9 ± 3.5 ed        | 298.9 ± 23.8 bc  | 1.64 ± 0.3 ab  | 27.5 ± 7.0 a-c        | 83.2 ± 9.8 bc             | 110.7 ± 5.2 b             |
| 20 Gy     | 241.7 ± 21.1 ab        | 131.2 ± 9.3 b-d       | 373.0 ± 29.5 b   | 1.83 ± 0.1 a  | 13.5 ± 3.5 bc         | 94.9 ± 6.6 b              | 108.4 ± 6.2 b             |

*Values are means ± SE, different letters show significant differences using Duncan’s test ($P$ value < 0.05)

### Table 5 Influence of chemical and physical agents on the content of total polyphenols (TPC), flavonoids, flavonols, anthocyanins, and tannins in *L. spectabilis* ‘Valentine’ after 11 weeks of in vitro culture

| Treatment   | TPC (mg g$^{-1}$) | Flavonoids (mg g$^{-1}$) | Flavonols (mg g$^{-1}$) | Anthocyanins (µg g$^{-1}$) | Tannins (µg g$^{-1}$) |
|-------------|-------------------|--------------------------|-------------------------|---------------------------|------------------------|
| Control     | 5.46 ± 0.11 c*    | 2.54 ± 0.13 a            | 1.33 ± 0.09 a           | 121.9 ± 5.0 ab            | 3.75 ± 0.04 c         |
| Gold nanoparticles  
50 ppm  | 6.68 ± 0.23 ab    | 2.39 ± 0.03 a            | 1.22 ± 0.02 a           | 118.3 ± 4.3 ab            | 4.01 ± 0.06 ab        |
| 75 ppm  | 6.62 ± 0.24 ab    | 2.44 ± 0.07 a            | 1.22 ± 0.03 a           | 129.7 ± 2.5 a             | 4.00 ± 0.06 ab        |
| 100 ppm | 6.23 ± 0.18 ab    | 2.57 ± 0.12 a            | 1.32 ± 0.02 a           | 113.9 ± 1.9 a-c           | 4.03 ± 0.02 a         |
| Microwaves  
9 s       | 6.48 ± 0.21 ab    | 2.48 ± 0.10 a            | 1.28 ± 0.10 a           | 120.4 ± 9.1 ab            | 3.87 ± 0.02 b-e       |
| 18 s       | 6.53 ± 0.39 ab    | 2.56 ± 0.08 a            | 1.39 ± 0.07 a           | 118.6 ± 2.0 a             | 3.87 ± 0.05 b-e       |
| 27 s       | 6.41 ± 0.04 ab    | 2.32 ± 0.07 a            | 1.22 ± 0.02 a           | 114.6 ± 4.2 a-c           | 3.93 ± 0.01 a-d       |
| X-rays     
10 Gy     | 6.96 ± 0.38 a     | 2.37 ± 0.10 a            | 1.21 ± 0.07 a           | 127.6 ± 1.6 ab            | 3.98 ± 0.05 a-c       |
| 15 Gy     | 8.86 ± 0.08 ab    | 2.47 ± 0.09 a            | 1.35 ± 0.05 a           | 106.5 ± 9.8 bc            | 3.84 ± 0.03 de        |
| 20 Gy     | 6.14 ± 0.15 b     | 2.41 ± 0.09 a            | 1.40 ± 0.03 a           | 97.6 ± 4.2 c              | 3.85 ± 0.05 c-e       |

*Values are means ± SE, different letters show significant differences using Duncan’s test ($P$ value < 0.05)
after 18 s microwave treatment and in the control bleeding hearts in the case of carotenes (Table 4). On the other hand, there was no impact of the tested factors on the ratio of total chlorophyll to carotenoids (data not shown).

All shoots produced from explants treated with AuNPs or electromagnetic irradiation contained more polyphenols (6.14–6.96 mg g\(^{-1}\) FW) than the untreated control (5.46 mg g\(^{-1}\) FW; Table 5). In contrast, there was no influence of the tested agents on the content of flavonoids (2.32–2.57 mg g\(^{-1}\) FW) and flavonols in bleeding heart (1.22–1.40 mg g\(^{-1}\) FW). None of the analyzed shoots contained more anthocyanins than the control, however, bleeding hearts irradiated with 20 Gy of X-rays contained markedly less pigments of this group (Table 5), resulting in an evident phenotype change (Fig. 1F). As for tannins, an increase in their concentration was detected in the experimental objects treated with AuNPs, regardless of nanoparticles concentration, and after treatment with the highest and lowest dose of microwaves and X-rays, respectively (Table 5).

**Estimation of nuclear DNA content in in vitro-derived plantlets by flow cytometry**

The genome size of *L. spectabilis* was found to be very small; approximately 1281 Mbp (1.268–1.351 pg/2C; mean 1.314 pg/2C). No changes in the ploidy level were reported in any of the 150 plants analyzed, regardless of treatment. However, a total of four plants (from 50 ppm AuNPs, 27 s MW, 10 Gy, and 20 Gy X-rays treatments) contained a markedly lower DNA content than the remaining specimens from those experimental combinations. Moreover, one specimen after 10 Gy X-irradiation contained significantly more DNA in its nuclei (Fig. 2). Plants treated with 20 Gy X-rays had the highest range of nuclear DNA content.

**In vitro rooting, acclimatization, and ex vitro growth**

All shoots regenerated adventitious roots on the IAA-supplemented rooting medium. The longest roots (6.0 cm), significantly different from the control (3.8 cm), were found in the experimental object with 100 ppm AuNPs in the culture medium (Table 6). On the other hand, plants treated with 15 and 20 Gy regenerated the shortest root systems (1.7–1.9 cm). None of the agents tested increased the area of the root system, but plants produced after X-irradiation, regardless of the dose, had a smaller root area (0.40–0.56 cm\(^2\)) and volume (6.7–10.5 mm\(^3\)) compared to the control (0.91 cm\(^2\); 18.9 mm\(^3\)). On the other hand, the application of 100 ppm AuNPs resulted in the smallest root diameter (Table 6). The highest (22.4) and lowest (5.3) numbers of forks were found in the root systems of plants treated with 100 ppm AuNPs and 20 Gy of X-rays, respectively. There was no difference in the number of tips between the untreated control and any other experimental object (Table 6).

A complete 100% survival of shoots was reported after the 2-week acclimatization, regardless of treatment (Fig. 1G). However, in the following weeks, some of the
plants eventually died. Plantlets from the control object and after 27 s treatment with microwaves had significantly lower survivability (86–88%) after 6 weeks of ex vitro growth than those from the remaining experimental combinations (96–100%) (Fig. 3A).

Over 80% of acclimatized plants treated with 75 ppm AuNPs, 18 s of microwaves, or 20 Gy of X-rays were of superb quality, with several new leaves developed. Only in the control object, the share of plants of the highest quality was below 50% (Fig. 3B). Most of the bleeding hearts did not differ phenotypically from the control (Fig. 1G). However, four plants (one after 75 ppm AuNPs treatment, two after 10 Gy X-rays irradiation, and one from the 20 Gy X-rays combination) were markedly different with an altered, more irregular shape of leaves (Fig. 1H).

**Analysis of genetic fidelity**

A total of 12,394 scorable bands were detected by five DAMD, four ISSR, five RAPD, and five SCoT primers in 80 plants (Table 7). The size of bands ranged from 152.9 to 2976.8 bp. One ISSR primer (I-3) failed to generated unambiguous amplicons; therefore, it was not included in the study. Among the molecular marker systems tested, DAMDs generated most polymorphisms: a total of 52.5% loci were polymorphic in six specimens (i.e. 7.5% of all plants analyzed). Primers D-1 and D-3 generated the highest number of bands (17 per sample). The latter one was also the most effective in screening for variation (six distinguished genotypes). On the other hand, primer D-4 produced only six amplicons per sample. This was also the only DAMD primer that did not detect polymorphisms in the present study. Specific loci were the least numerous (2.0 per primer), followed by polymorphic (4.4) and monomorphic ones (5.8). As for RAPD analysis, three primers (R-3, R-4, and R-5) detected polymorphic and specific bands in five plants (6.25% of all plants tested). Similarly, four SCoT primers (all except for S-1) generated polymorphic amplicons in five specimens. No polymorphisms were detected by ISSRs. This marker system detected also the lowest mean number of bands (480 per primer). An example visualization of PCR products is shown in Fig. 4. No polymorphisms were found in the control plants.

Among the three marker systems, which detected polymorphic bands, the highest mean H, PIC, and MI values were reported with DAMDs (Table 8). Those markers also had the highest mean discriminating power (0.492), followed by RAPD (0.402) and SCoT (0.286). On the other hand, the highest resolving power was found with RAPDs (0.360). According to the PCoA analysis, three genotypes obtained as a result of treatment with 20 Gy X-rays (two genotypes) and 100 ppm AuNPs (one genotype) were the most distinct from the predominant control, based on the data provided by three marker systems (Fig. 5). The genotype obtained after 75 ppm AuNPs treatment was also distant from the control according to RAPD and SCoT markers, although not so much according to DAMDs. Similarly, various positioning of the genotype created after MW irradiation for 27 s was found with DAMD and SCoT markers. A relatively small distance between the control group and genotype obtained after 15 Gy X-rays treatment was detected by DAMDs.

The AMOVA analysis; based on DAMD, RAPD, and SCoT markers; confirmed a significant influence of the experimental treatments on the occurrence of interspecific genetic variation in bleeding heart ‘Valentine’ (data not shown). The comparison of the efficacy of various marker

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**Table 6** Influence of chemical and physical agents on the root system architecture in *L. spectabilis* ‘Valentine’ after 6 weeks of in vitro rooting culture on the MS medium with 2.0 mg L⁻¹ IAA

| Treatment         | Total length (cm) | Area (cm²) | Volume (mm³) | Diameter (mm) | No. of forks | No. of tips |
|-------------------|-------------------|------------|--------------|---------------|--------------|-------------|
| Control           | 3.8 ± 0.6 bc*     | 0.91 ± 0.11 a | 18.9 ± 1.3 a | 0.87 ± 0.05 a | 13.1 ± 2.3 b-d | 18.0 ± 4.7 a-d |
| Gold nanoparticles|                   |            |              |               |              |             |
| 50 ppm            | 4.6 ± 0.5 ab      | 0.89 ± 0.08 a | 14.6 ± 1.3 ab | 0.69 ± 0.02 bc | 17.5 ± 2.9 ab | 25.5 ± 1.6 ab |
| 75 ppm            | 4.2 ± 0.8 bc      | 0.90 ± 0.11 a | 16.2 ± 1.6 a  | 0.77 ± 0.07 ab | 15.9 ± 2.6 a-c | 20.9 ± 3.2 a-c |
| 100 ppm           | 6.0 ± 0.9 a       | 1.04 ± 0.10 a | 14.9 ± 1.5 ab | 0.59 ± 0.04 c  | 22.4 ± 3.4 a  | 27.1 ± 4.1 a  |
| Microwaves        |                   |            |              |               |              |             |
| 9 s               | 3.6 ± 0.5 b-d     | 0.77 ± 0.07 ab | 13.7 ± 1.0 a-c | 0.72 ± 0.05 a-c | 13.0 ± 1.7 b-d | 16.5 ± 2.9 b-d |
| 18 s              | 3.0 ± 0.6 b-e     | 0.77 ± 0.13 ab | 16.9 ± 2.7 a  | 0.89 ± 0.03 a  | 11.1 ± 2.2 b-e | 11.4 ± 2.4 cd |
| 27 s              | 3.4 ± 0.6 b-e     | 0.79 ± 0.11 ab | 16.3 ± 2.2 a  | 0.86 ± 0.08 ab | 11.3 ± 1.8 b-e | 15.5 ± 2.2 b-d |
| X-rays            |                   |            |              |               |              |             |
| 10 Gy             | 2.5 ± 0.3 c-e     | 0.56 ± 0.06 bc | 10.5 ± 1.2 b-d | 0.76 ± 0.05 a-c | 9.8 ± 2.2 c-e | 14.9 ± 2.9 cd |
| 15 Gy             | 1.9 ± 0.5 de      | 0.40 ± 0.09 c  | 6.7 ± 1.1 e   | 0.75 ± 0.07 a-c | 5.8 ± 2.1 de  | 14.7 ± 5.1 cd |
| 20 Gy             | 1.7 ± 0.2 e       | 0.44 ± 0.06 c  | 9.1 ± 1.4 de  | 0.88 ± 0.03 a  | 5.3 ± 0.6 e  | 8.9 ± 1.8 d  |

*Values are means ± SE, different letters show significant differences using Duncan’s test (P value < 0.05)*
types is shown in Table 9. Noticeably, all plants which showed phenotype alternations were also distinguished by SPAR and/or FCM analyses (mutants M2, M5, M6, M8). However, five genotypes that were distinguished by (cyto) genetic markers did not have evident phenotype changes.

Discussion
The development of efficient in vitro culture systems is imperative for the large-scale production, acquisition
Table 7 Comparative analysis of molecular products obtained from L. spectabilis ‘Valentine’ plants analyzed with DAMD, ISSR, RAPD, and SCoT markers

| Primer code | Primer sequence 5’ → 3’ | Total no. of bands | Size of bands (bp) | No. of loci | No. and (%) of plants with polymorphism | No. of genotypes |
|-------------|--------------------------|--------------------|-------------------|------------|----------------------------------------|-----------------|
|             |                          | Min               | Max               | Total      | Mono       | Poly       | Spec       |               |               |
| DAMD        |                          |                   |                   |            |            |            |            |               |               |
| D-1         | AGGACTCGATAAACAGGCTCC    | 1200              | 386.1             | 2350.5     | 17         | 13         | 2          | 2           | 1 (1.25)     | 2              |
| D-2         | CCCAGCAACTGATCGCACAC     | 247               | 416.0             | 1008.4     | 8          | 1          | 4          | 3           | 4 (5.0)      | 4              |
| D-3         | AAATGGGGCAACGTCTGGT      | 655               | 364.7             | 338.7      | 17         | 7          | 6          | 4           | 5 (6.25)     | 6              |
| D-4         | GTGTGCACTAGTGCTGGG       | 480               | 514.6             | 1472.9     | 6          | 6          | 0          | 0           | 0 (0.0)      | 1              |
| D-5         | ACATCGCAAGTGACACAGG      | 641               | 666.6             | 2890.7     | 13         | 2          | 10         | 1           | 4 (5.0)      | 4              |
| **Σ**       |                          | **3223**          | –                 | –          | **61**     | **29**     | **22**     | **10**      | **6 (7.5)**  | **7**          |
| (mean)      |                          | (644.6)           |                   |            |            |            |            |             |               |                |
| ISSR        |                          |                   |                   |            |            |            |            |             |               |                |
| I-1         | GAG GGT (GGA)_2 TCT      | 320               | 296.2             | 1318.1     | 4          | 4          | 0          | 0           | 0 (0.0)      | 1              |
| I-2         | C (GAGA)_4               | 400               | 755.6             | 2185.2     | 5          | 5          | 0          | 0           | 0 (0.0)      | 1              |
| I-3         | C (AGAG)_4               | –                 | –                 | –          | –          | –          | –          | –           | –              |                |
| I-4         | (GACA)_4                 | 800               | 407.0             | 1461.4     | 10         | 10         | 0          | 0           | 0 (0.0)      | 1              |
| I-5         | (GGA)_6                  | 400               | 424.1             | 870.1      | 5          | 5          | 0          | 0           | 0 (0.0)      | 1              |
| **Σ**       |                          | **1920**          | –                 | –          | **24**     | **24**     | **0**      | **0**       | **0 (0.0)**  | **1**          |
| (mean)      |                          | (480.0)           |                   |            |            |            |            |             |               |                |
| RAPD        |                          |                   |                   |            |            |            |            |             |               |                |
| R-1         | GCT GCC TCA GG           | 800               | 152.9             | 1452.2     | 10         | 10         | 0          | 0           | 0 (0.0)      | 1              |
| R-2         | CAA TCG CGG T            | 720               | 220.1             | 2823.0     | 9          | 9          | 0          | 0           | 0 (0.0)      | 1              |
| R-3         | GGT GAC GCA G            | 817               | 335.4             | 2749.3     | 23         | 4          | 16         | 3           | 5 (6.25)     | 5              |
| R-4         | CCC AGT CAC T            | 573               | 241.2             | 2281.9     | 14         | 7          | 4          | 3           | 4 (5.0)      | 4              |
| R-5         | GGT GAC GCA G            | 642               | 322.8             | 1110.9     | 11         | 6          | 4          | 1           | 4 (5.0)      | 3              |
| **Σ**       |                          | **3552** (710.4)  | –                 | –          | **67**     | **36**     | **24**     | **7**       | **5 (6.25)** | **6**          |
| (mean)      |                          | (480.0)           |                   |            |            |            |            |             |               |                |
| SCoT        |                          |                   |                   |            |            |            |            |             |               |                |
| S-1         | CAA CAA TGG CTA CCA CCG  | 720               | 546.2             | 2768.6     | 9          | 9          | 0          | 0           | 0 (0.0)      | 1              |
| S-2         | CAA CAA TGG CTA CCA CCT  | 641               | 335.4             | 2288.4     | 9          | 8          | 0          | 1           | 1 (1.25)     | 2              |
| S-3         | CAA CAA TGG CTA CCA CGT  | 876               | 345.0             | 2976.8     | 13         | 5          | 6          | 2           | 1 (1.25)     | 2              |
| S-4         | ACG ACA TGG CGA CCA AC   | 650               | 597.0             | 1793.6     | 12         | 8          | 3          | 1           | 4 (5.0)      | 3              |
| S-5         | ACG ACA TGG CGA CCA TCG  | 812               | 209.0             | 1838.5     | 13         | 10         | 3          | 0           | 5 (6.25)     | 2              |
| **Σ**       |                          | **3699**          | –                 | –          | **56**     | **40**     | **12**     | **4**       | **5 (6.25)** | **4**          |
| (mean)      |                          | (739.8)           |                   |            |            |            |            |             |               |                |

*DAMD* directed amplification of minisatellite DNA, *ISSR* inter sequence simple repeat, *RAPD* randomly amplified polymorphic DNA, *SCoT* start codon target polymorphism, *mono.* monomorphic, *poly.* polymorphic, *spec.* specific (unique; present in a single band profile)
Fig. 4 Example DAMD band profiles of *L. spectabilis* ‘Valentine’ received as a result of electrophoresis of the DNA amplification products obtained with primer D-1. Outermost lanes (wm) are DNA bp weight markers, while inner lines represent plants treated with gold nanoparticles (AuNPs), microwaves (MW), X-rays, and the non-treated control. Arrows indicate polymorphic genotypes.

Table 8 Values of Heterozygosity index (H), Polymorphic Information Content (PIC), Effective multiplex ratio (E), Marker Index (MI), Discriminating power (D), and Resolving power (R) of the marker systems used in the study.

| Primer | H    | PIC  | E     | MI    | D    | R    |
|--------|------|------|-------|-------|------|------|
| DAMD   |      |      |       |       |      |      |
| D-1    | 0.208| 0.186| 15.00 | 0.0023| 0.222| 0.100|
| D-2    | 0.474| 0.362| 3.088 | 0.0023| 0.851| 0.275|
| D-3    | 0.499| 0.375| 8.188 | 0.0030| 0.768| 0.475|
| D-4    | 0    | 0    | 0     | 0     | 0    | 0    |
| D-5    | 0.473| 0.361| 8.012 | 0.0036| 0.620| 0.625|
| Mean   | 0.331| 0.257| 6.858 | 0.0022| 0.492| 0.295|
| ISSR   |      |      |       |       |      |      |
| Mean   | 0    | 0    | 0     | 0     | 0    | 0    |
| RAPD   |      |      |       |       |      |      |
| R-1    | 0    | 0    | 0     | 0     | 0    | 0    |
| R-2    | 0.494| 0.372| 10.21 | 0.003 | 0.803| 1.125|
| R-3    | 0.500| 0.375| 7.163 | 0.003 | 0.738| 0.325|
| R-4    | 0.395| 0.317| 8.025 | 0.004 | 0.468| 0.350|
| R-5    | 0.278| 0.213| 5.080 | 0.002 | 0.402| 0.360|
| Mean   | 0    | 0    | 0     | 0     | 0    | 0    |
| SCoT   |      |      |       |       |      |      |
| S-1    | 0    | 0    | 0     | 0     | 0    | 0    |
| S-2    | 0.195| 0.176| 8.013 | 0.002 | 0.208| 0.025|
| S-3    | 0.266| 0.230| 10.95 | 0.003 | 0.291| 0.200|
| S-4    | 0.437| 0.342| 8.125 | 0.004 | 0.542| 0.250|
| S-5    | 0.342| 0.284| 10.15 | 0.003 | 0.391| 0.300|
| Mean   | 0.248| 0.206| 7.448 | 0.002 | 0.286| 0.155|

*DAMD* directed amplification of minisatellite DNA, *ISSR* inter sequence simple repeat, *RAPD* randomly amplified polymorphic DNA, *SCoT* start codon target polymorphism.
of secondary metabolites, and breeding possibilities in ornamental and medicinal plant species. This is the first study on the application of nanoparticles and ionizing radiation in micropropagation and mutation induction in bleeding heart. The results of the current experiment have broadened the scope of knowledge about the biology and requirements of *L. spectabilis*, which is a rather difficult plant species to be grown in the in vitro environment (VITROFLORA, Trzęsacz, Poland, personal communication).

### In vitro development and acclimatization efficiency of bleeding heart exposed to AuNPs and electromagnetic irradiation

The present study highlights the utility of gold nanoparticles in micropropagation of bleeding heart ‘Valentine’. The addition of AuNPs into the culture medium, regardless of their concentration, stimulated the proliferation of shoots and nearly tripled the propagation ratio (at 75 ppm).
compared to the non-treated control. Moreover, AuNPs at the highest 100 ppm concentration enhanced the elongation and branching of roots during the rooting stage. This is of significant importance for the whole plant life as roots provide and transport water and minerals to the upper organs that permit successful photosynthetic nutrition (Dubrovsky et al. 2019). Similarly to our findings, the best effects in the in vitro stimulation of the rosulate and phyllomorphs regeneration in *Streptocarpus × hybridus* Voss. were reported if silver or gold nanoparticles (20 nm in size) were added to the MS medium but at a much lower concentration of 10 ppm (Tymoszuk and Miler 2019). According to Tian et al. (2018), the application of 25 ppm of manganese nanoparticles (30 nm) can result in pores formation in the roots stimulating a greater uptake of water and nutrients and, consequently, in significant plant growth. *Allium cepa* L. root tip cells treated with AuNPs exhibited an increment in the frequency of mitotic index with the increase in nanoparticles concentration from 10 to 1000 ppm (Gopinath et al. 2013). Moreover, nanoparticles can additionally alter the absorption of metal ions important for plant development (Timoteo et al. 2019) or stimulate a more intensive lignin accumulation in cell walls improving the quality of in vitro propagated plants (Bernard et al. 2015). These mechanisms of NPs action could explain the present results and the activation of axillary meristems in bleeding heart. Nonetheless, less beneficial effects of various types of nanoparticles in other plant species are also known, such as inhibition of seed germination and/or seedling development (Sanzari et al. 2019). Further studies should focus on the effect of NPs size on the micropropagation efficiency in bleeding heart.

In the current study, the foliar analysis revealed that treatment with X-rays, especially with 15 and 20 Gy, increased the fresh and dry weight of shoots without altering their relative water content. Those plants were also of the highest viability when transferred to ex vitro conditions (100% survival). This positive effect of X-rays could be related to the so-called hormesis, i.e. a favorable response of plants to lower doses, which is especially important if the studied factors were used at lower doses, which is especially important for AuNPs application in micropropagation.

### Physiological and biochemical response of bleeding heart to in vitro-induced stress

The present research confirmed that AuNPs, MW, and X-rays are a source of considerable oxidative stress in plants if applied at high doses. This was evident based on the activity analysis of the three enzymes studied, which was elevated up to five times as in the case of G-POD after applying 20 Gy of X-rays. Our results are in agreement with several other studies related to the antioxidant defense system in higher plants (Barbacz et al. 2016; Chandel et al. 2017). Nanoparticles and electromagnetic radiation have various chemical effects, including causing deterioration in large molecules in cells and imbalance in ionic equilibrium leading to generation of hazardous by-products, known as reactive oxygen species (ROS), during biological reactions (Dietz and Herth 2011; Kivrak et al. 2017). In the present study, guaiacol peroxidase was the most sensitive marker of oxidative stress, while glutathione reductase—the least (a twofold activity increase reported solely in the experimental object with the highest concentration of AuNPs). These findings corroborate our previous report on the impact of cryopreservation stress on the physiological response in *L. spectabilis* (Kulus and Tymoszuk 2021). On the other hand, the antioxidant enzymatic activity did not change significantly if the studied factors were used at lower doses, which is especially important for AuNPs application in micropropagation.

It was also found that both chemical and physical agents can be used to alternate the profile of metabolites in bleeding heart. Spectral fingerprinting revealed modulation of plant pigments; a significant, nearly two-fold increase in the concentrations of chlorophylls and carotenoids were found if microwaves were used for a total of 18 s. Other changes were also found that lead to the plantlets color change during the in vitro growth stage. According to Kumar et al. (2018), plants produce a number of secondary metabolites such as flavonoids, phenolics, terpenoids, amino acids and increase the level of different enzymes to reduce the effect of ROS, which is the probable reason for those changes. It is worth...
mentioning that all experimental objects contained more polyphenols (TPC) than the control and, often also, a higher concentration of tannins, which is of significant importance for the pharmaceutical industry. Gold nanoparticles in particular affected the content of tannins, which additionally highlights the utility of NPs in plant biotechnology and abiotic elicitation (Halder et al. 2019).

Genome size and genetic stability analyses in bleeding heart

Currently, the genome size (i.e. the total amount of DNA contained in the cell nucleus) is estimated in over 12,000 plant species (Pellicer and Leitch 2020). This parameter can have an impact at many scales, from influencing gene and genome dynamics to playing a role at the whole-plant level, influencing, for example, plant growth strategies (Dodsworth et al. 2015; Guignard et al. 2016). The obtained results revealed that bleeding heart has a very small genome size of approximately 1281 Mbp (2C DNA = 1.314 pg), according to the classification of angiosperms by Soltis et al. (2003). In comparison, the lowest known value of this parameter in Genlisea tuberosa Rivadavia (Lentibulariaceae) was 61 Mbp/1C, and the largest in Paris japonica (Franch. & Sav.) Franch (Liliaceae) reached 149,000 Mbp/1C of DNA (Pellicer et al. 2018). A plant species of such a small genome may be a good model organism in experimental biology, for example in the studies related to understanding evolutionary trajectories and ecosystem dynamics.

In the present study, a total of five plants (3.3%) had a changed DNA content, mainly after irradiation with X-rays (three plants). Most of the altered bleeding hearts (80%) had a lower nuclear DNA concentration than the remaining plants from the same experimental combination. This suggests that the applied mutagens caused considerable deletions from the genome of the original ‘Valentine’ cultivar. This also may be a result of stress, which negatively affects the synthesis of DNA, as reported in other plant species (Oladosu et al. 2016). A similar negative effect of ionizing irradiation on DNA content in chrysanthemum was reported previously by Miler et al. (2020). In bleeding heart, only in a single sample, treated with 10 Gy X-rays, an increase in genome size was found.

Since some genetic alternations can be eliminated due to the so-called diplontic selection and mutational meltdown (Klekowski 2003), the genetic (PCR) analysis of plants was performed after 6 weeks of ex vitro growth. The SPAR markers detected mutations in six of the analyzed plants (7.5%). Electromagnetic/ionizing radiation is well known for its mutagenic character and is broadly used in classical mutation breeding (Miler and Kulus 2018; Miler et al. 2021). Nanoparticles also can penetrate cells and impair stages of mitosis causing chromatin bridges and stickiness, disturbed metaphase, multiple chromosomal breaks, and cell disintegration (Kumari et al. 2009). To date, the mutagenic character of zinc oxide nanoparticles (ZnONPs) and AgNPs was confirmed by Kumari et al. (2011) in onion and by Tymoshuk and Kulus (2020) in chrysanthemum, respectively. The present study confirmed also the genotoxic effect of AuNPs if applied at high concentrations. Based on the frequency of variation occurrence in L. spectabilis, the usefulness of the three studied factors in breeding can be arranged as follows: MW < AuNPs < X-rays.

In the present study, only in a single plant (M8; X-ray treated), all three marker types (FCM, SPAR, and phenotype analysis) detected variation, while in the mutant M2 (AuNPs-treated) changes in the DNA sequence and plant morphology were evident. Lack of noticeable variation in the DNA content in the latter plant could result from the small size of the mutation, as most of the DNA sequence alternations between alleles in plants are limited to single nucleotides (Morgil et al. 2020). On the other hand, in the mutant M1, only FCM detected a significant change in the nuclear DNA amount. As for this plant, the variation could have occurred in the part of the genome, which is not covered by SPAR markers (Nadeem et al. 2017). The same explanation goes for mutants M5 and M6 (Table 9). In the remaining four mutants, genetic and/or cytogenetic markers detected variation despite no phenotype changes. This may suggest that the DNA was altered in the non-coding regions of the genome; the so-called gene deserts or ‘junk DNA’ (Nadeem et al. 2017). All of the phenotype alternations reported in the present study considered changes in the shape and symmetry of leaves. This is one of the most common changes occurring as a result of mutation induction (Masters 1869), important in bleeding heart as the species is also valued for its distinctive foliage with cut merit (Hodges 2012). Forthcoming analyses will be aimed at analyzing the parameters of flowers in blooming plants.

The present research confirmed the diverse usefulness of various PCR-based marker systems in the detection of polymorphisms in bleeding heart. Among the four marker systems tested, only ISSR failed to detect polymorphic bands. Moreover, this was the only marker where not all of the used primers generated amplicons. Similarly, in Vitis vinifera L., ten ISSR primers were evaluated, but only two were capable to amplify distinct and reproducible bands with sizes between 300 and 2000 bp (Lazo Javalera et al. 2016). The discriminating power of the studied markers could be arranged in the following order: ISRR < SCOT < RAPD < DAMD. DAMDs distinguished the highest number of polymorphic plants, although the PCR reaction for those markers was the most difficult to optimize (data not shown). The high efficiency of RAPDs was also confirmed in members of the Asteraceae botanical family by Lema-Rumińska et al. (2019).
In future studies, other cultivars of bleeding heart should be tested to obtain the widest possible variability, as it is known that the genetic factor has a significant impact on the efficiency and range of spontaneous and induced mutations occurrence. For example, in chrysanthemum, cultivars with purple inflorescences mutate most often, while the least frequent variation occurs in plants with yellow flowers (Schum 2003). For this reason, also in the present study, a red-floral ‘Valentine’ cultivar was used, but the purple-pink L. spectabilis ‘Gold Heart’ could also provide some interesting results.

It is also worth mentioning that in the present study, none of the experimental objects differed significantly from the control in terms of viability during the in vitro cultivation (86–100%). For nodal segments of Citrus macrophylla Wester., the LD₅₀ was determined as 29.2 Gy of γ-rays (Tallón et al. 2015). This underlines the high tolerance of bleeding heart to stress conditions. It is, therefore, possible that this species could be a source of resistance genes in hybridization programs. In future studies with bleeding heart, an even higher dose of the tested factors could be used to increase the number of mutated plants and traits and to determine the LD₅₀ in this species.

Conclusions

The addition of AuNPs into the culture medium, especially at 75 ppm, provokes the proliferation of shoots and multiplication ratio in Lamprocapnos spectabilis ‘Valentine’. Gold nanoparticles also stimulate the elongation and branching of roots. On the other hand, irradiation of explants with X-rays results in the development of leaves with the highest fresh and dry weight but inhibits the rooting of shoots. Gold nanoparticles and electromagnetic irradiation affect also the profile of metabolites in bleeding heart and improve the survival and quality of acclimatized plants. Both chemical and physical agents applied at high doses can be a source of genetic variation, with X-rays and MW being the most and the least effective mutagens, respectively. DAMD and RAPD markers are the most suitable for mutation detection. The induced biochemical and genetic alternations can result in phenotype change in L. spectabilis leading to the creation of new cultivars. The simple propagation and breeding layouts could be easily implemented in commercial and research laboratories with possible extensions to other cultivars or species.

Future studies should focus on the usage of other agents: such as gamma rays, fast neutrons (FN), or AgNPs, applied at even higher doses than in the current study, in mutagenesis of bleeding heart. Various cultivars should be compared to screen for traits indicating higher susceptibility to mutation occurrence. A broad analysis of phenotype alterations should be performed in the in vivo-grown plants to confirm their stability in the subsequent generations.

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Declarations

Conflict of interest The authors declare no commercial or financial conflict of interest.

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