

Mutagenic Effectivity of Cadmium Sulphide and Copper Oxide Nanoparticles on Some Physiological and Cytological Attributes of *Lathyrus sativus* L.

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Received July 5, 2016; accepted March 5, 2017

Summary Dry seeds (moisture content: 17.50%) of *Lathyrus sativus* L. (Family: Fabaceae, common name—grass pea) are exposed (doses: 0.25, 0.50 and 1.00 $\mu\text{g mL}^{-1}$, duration: 3 and 6 h) to chemically synthesized cadmium sulphide and copper oxide nanoparticle (NP) treatments for assessment of genotoxic potentiality. The objective of the work is to foresee whether the NPs possess effective mutagenic potentiality. Result is significant in the direction of underlined objective.

Key words *Lathyrus sativus*, CdS-NP, CuO-NP, Physiological and cytological attributes, Mutagenic potentiality.

Nanoparticles (NPs; size <100 nm) possess significant uses in industry, biomedical sciences and electronic devices (Manickathai *et al.* 2008, Masarovičová and Kráľová 2013), plant biotechnology and agriculture (Scrinis and Lyons 2007) including crop protection (Nair *et al.* 2010). NP-mediated plant interaction is opening a new dimension of research on induced mutagenesis (Halder *et al.* 2015a, 2015b, Kumbhakar *et al.* 2016). Doses of NPs administered and species level sensitivity are significant pre-requisites for designing experiments on mutation breeding. With the view to it, the present investigation describes the effective potentiality of chemically synthesized CdS and CuO-NPs (semiconductor NPs; band gap—CdS: 2.45 eV, CuO: 3.149 eV) on some physiological (Petri plate seed germination and seedling length—for the assessment of phytotoxicity) and cytological (mitotic aberrations—for determination of genotoxicity) attributes of *Lathyrus sativus* L. (Family: Fabaceae; common name—grass pea, a legume of commercial importance).

Materials and methods

Germplasm

Seeds (moisture content, 17.50%; length, 4.94 ± 0.09 mm; breadth, 4.40 ± 0.13 mm) of *L. sativus* were obtained from Horticultural Research Station, Director-

ate of Principle Agriculture Officer, Nadia, Govt. of West Bengal, India.

Preparation of NPs

CdS-NPs were prepared following wet chemical co-precipitation techniques adopted by Halder *et al.* (2015a). CuO-NPs were chemically synthesized according to Topnani *et al.* (2010). Bulk compounds were prepared without the application of capping agents.

Both the NPs were characterized based on different opto-physical parameters (data unpublished). Instrumentation techniques revealed corroborating result of the size and shape for both the nano-crystals and they were with conformation of the nano-standard quality.

Treatments

Dry seeds of *L. sativus* were treated with CdS and CuO-NPs (doses: 0.25, 0.50 and 1.00 $\mu\text{g mL}^{-1}$; duration: 3 and 6 h). Controls (dry: bulk CdS—0.25 $\mu\text{g mL}^{-1}$, 3 h and bulk CuO—0.25 $\mu\text{g mL}^{-1}$, 3 h) were also kept for assessment. A hundred seeds were exposed in each lot of treatments.

Seed germination and seedling growth

Treated and control seeds (100 in each case) were given in Petri plates lined with moist filter paper (28 ± 1 °C). Emergence of radicle was considered as index of germination. Twenty randomly selected seedlings from each set were measured on the seventh day from treatments in a millimeter graph paper. Lethality and injury were

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DOI: 10.1508/cytologia.82.267

calculated from germination frequency and seedling length, respectively (Konzak *et al.* 1965), and represented as percent of controls (bulk CdS and CuO).

Mitotic study

Germinating roots of about 2 mm in length from each set (three to four) were cut at uniform duration (12 noon to 12:30 p.m.), fixed in 1:3 (v/v) acetic alcohol overnight and stored in 70% alcohol in a refrigerator ($16\pm 1^\circ\text{C}$). Roots were stained in orcein-HCl (9:1) mixture for 3 to 4 h following standard hydrolysis procedure. Root tips were squashed in 45% acetic acid and observed under an Olympus trinocular light microscope (Magnus MLX-DX).

Mitotic index [(total number of dividing cells/total cells scored) $\times 100$], types of aberration (both in dividing and resting cells) and total abnormality percentage in dividing cell were assessed. Photomicrographs were taken from suitable squash preparations.

Results and discussion

Germination frequency and seedling growth

Results of seed germination frequency and seedling length are presented in Table 1. Inhibition in seed germination frequency is recorded in treatments (CdS-NPs: 92.0–76.0%; CuO-NPs: 90.0–72.0%) than controls (dry: 96.0%, bulk CdS: 94.0% and bulk CuO: 93.0%) and is marked in higher doses ($1.00\ \mu\text{g mL}^{-1}$, 6 h). Seedling length also reduces significantly ($p < 0.05$) in treatments (CdS-NPs: 24.82 ± 1.19 to 11.33 ± 0.61 mm; CuO-NPs: 17.28 ± 0.62 to 9.44 ± 0.46 mm) than controls. Maximum reduction is observed in $0.25\ \mu\text{g mL}^{-1}$, 3 h for CdS-NPs and $1.00\ \mu\text{g mL}^{-1}$, 3 h for CuO-NPs. On comparative

basis of unit doses of NPs, it seems that germination frequency is relatively more affected in CuO-NPs treatment, while seedling growth did not manifest any clear response.

Assessment of lethality indicates that none of the employed doses has shown 50.0% reduction in germination. It suggests that the doses of NPs administered are tolerant to the genotype. Injury ranges from 8.62 to 58.28% in CdS-NPs and 13.08 to 52.52% in CuO-NPs. The inhibitory action of NPs on the studied physiological attributes accentuates induction of phytotoxicity. NPs are reported to inhibit seed germination, seedling growth, and root elongation among others in different plant species (Lee *et al.* 2010, Ma *et al.* 2010, Atha *et al.* 2012, Shaymurat *et al.* 2012, Kumbhakar *et al.* 2016), thereby conferring phytotoxicity. However, enhancement in different physiological attributes has also been studied (Lu *et al.* 2002, Lin *et al.* 2004, Masarovičová and Kráľová 2013).

Reduction in germination frequency and seedling growth as the consequence of mutagen treatments have been attributed to the nature and the extent of chromosomal aberration occurring in the cells (Sax 1942, Lea 1946, Read 1959, Evans and Sparrow 1961, Datta and Biswas 1983) and also to structural changes (Gray and Read 1950).

Mitotic study

Compared to controls, mitotic index reduces significantly ($p < 0.05$) in treatments; while types and total aberration frequency estimated from dividing cells (show dose dependent relationship) enhances (Table 2) significantly ($p < 0.05$). NPs are therefore producing mitodepressive and genotoxic effects. Evans (1965) consid-

Table 1. Germination frequency and seedling length in controls and NP treatments.

Doses (% , $\mu\text{g mL}^{-1}$)	Duration (h)	Total no. of seeds given in Petri plate	Germination frequency (%)	Seedling length (mm)		Lethality*	Injury*
				Mean	SE		
Dry control		100	96.0	29.01	1.11	—	—
Bulk CdS control	3	100	94.0	27.16	0.92	—	—
CdS-NPs							
0.25	3	100	92.0	11.33	0.61	2.13	58.28
0.50	3	100	87.0	13.82	0.73	7.45	49.12
1.00	3	100	86.0	24.82	1.19	8.51	8.62
0.25	6	100	90.0	14.43	0.61	4.26	46.87
0.50	6	100	82.0	19.76	0.84	12.77	27.25
1.00	6	100	76.0	16.34	0.89	19.15	39.84
Bulk CuO control	3	100	93.0	19.88	0.70	—	—
CuO-NPs							
0.25	3	100	90.0	17.28	0.62	3.23	13.08
0.50	3	100	83.0	11.25	0.55	10.75	43.41
1.00	3	100	78.0	9.44	0.46	16.13	52.52
0.25	6	100	88.0	16.12	0.51	5.38	18.91
0.50	6	100	82.0	14.21	0.66	11.83	28.52
1.00	6	100	72.0	12.67	0.64	22.58	36.27
CD at 5% level				0.95			

* Percent of control.

Table 2. Mitotic index and frequency of aberration types in controls and NP treatments.

Mitotic index (%) and categorization of aberration types	Dry control	Bulk CdS control	CdS-NPs ($\mu\text{g mL}^{-1}$)						Bulk CuO control	CuO-NPs ($\mu\text{g mL}^{-1}$)						CD at 5% level
			3h			6h				3h			6h			
			0.25	0.50	1.00	0.25	0.50	1.00		0.25	0.50	1.00	0.25	0.50	1.00	
Mitotic index (%)	19.43	17.01	10.72	8.16	14.34	9.60	7.08	11.84	15.95	10.06	12.11	13.65	11.12	13.22	14.75	0.85
Physiological																
1. Stickiness	0.42	2.04	3.29	3.45	2.95	5.29	5.52	4.77	2.50	2.35	3.06	4.17	2.47	3.34	5.00	
2. Fragmentation																
a. Metaphase																
i. Gap	0.00	0.00	0.62	0.73	0.70	1.18	0.69	1.32	0.00	0.84	0.87	0.91	0.93	0.56	0.78	
ii. Fragments including paired fragments	0.00	0.00	1.03	0.73	0.98	0.88	0.92	1.15	0.00	0.67	0.73	1.09	0.62	0.84	1.25	
b. Anaphase																
i. Paired fragments	0.00	0.00	0.82	0.91	0.84	1.18	1.15	1.48	0.00	0.50	1.16	0.73	0.62	0.56	0.94	
c. Resting cells																
i. Micronuclei	0.00	0.00	0.00	0.13	0.21	0.00	0.18	0.27	0.00	0.00	0.24	0.49	0.17	0.32	0.51	
3. Inter-and intra-chromosomal rearrangements																
a. Metaphase																
i. Ring	0.00	0.00	0.00	0.00	0.28	0.00	0.23	0.82	0.00	0.00	0.29	0.73	0.00	0.97	1.25	
ii. Bridges	0.00	0.00	1.23	1.09	2.24	2.35	2.07	2.80	0.00	1.01	1.31	1.20	1.85	1.95	2.50	
b. Anaphase																
i. Pseudochiasma	0.00	0.00	2.26	2.36	1.68	2.35	2.53	2.14	0.00	1.68	1.46	2.54	1.55	1.25	2.34	
4. Spindle aberration																
a. Metaphase																
i. Polyploid	0.00	0.00	0.41	1.27	1.40	0.88	1.38	1.64	0.00	0.84	1.16	1.81	0.77	1.53	1.86	
b. Anaphase																
i. Laggards	0.00	0.00	1.03	1.27	1.82	2.06	2.07	1.97	0.19	0.67	1.60	2.18	1.39	1.67	2.19	
ii. Tripolarity	0.00	0.00	0.62	0.73	0.70	1.47	1.38	2.30	0.00	1.51	1.46	2.36	1.55	2.09		
6. Metabolic defects																
a. Giant cells	0.00	0.00	0.00	0.15	0.26	0.00	0.19	0.29	0.00	0.17	0.28	0.37	0.23	0.34	0.41	
b. Giant cell with vacuoles	0.00	0.00	0.00	0.00	0.05	0.00	0.00	0.07	0.00	0.00	0.00	0.06	0.00	0.04	0.16	
Total abnormality (%) in dividing cells	0.42	2.04	11.32	12.52	13.60	17.65	18.85	20.39	2.70	10.08	13.10	18.51	11.75	14.76	20.78	0.84

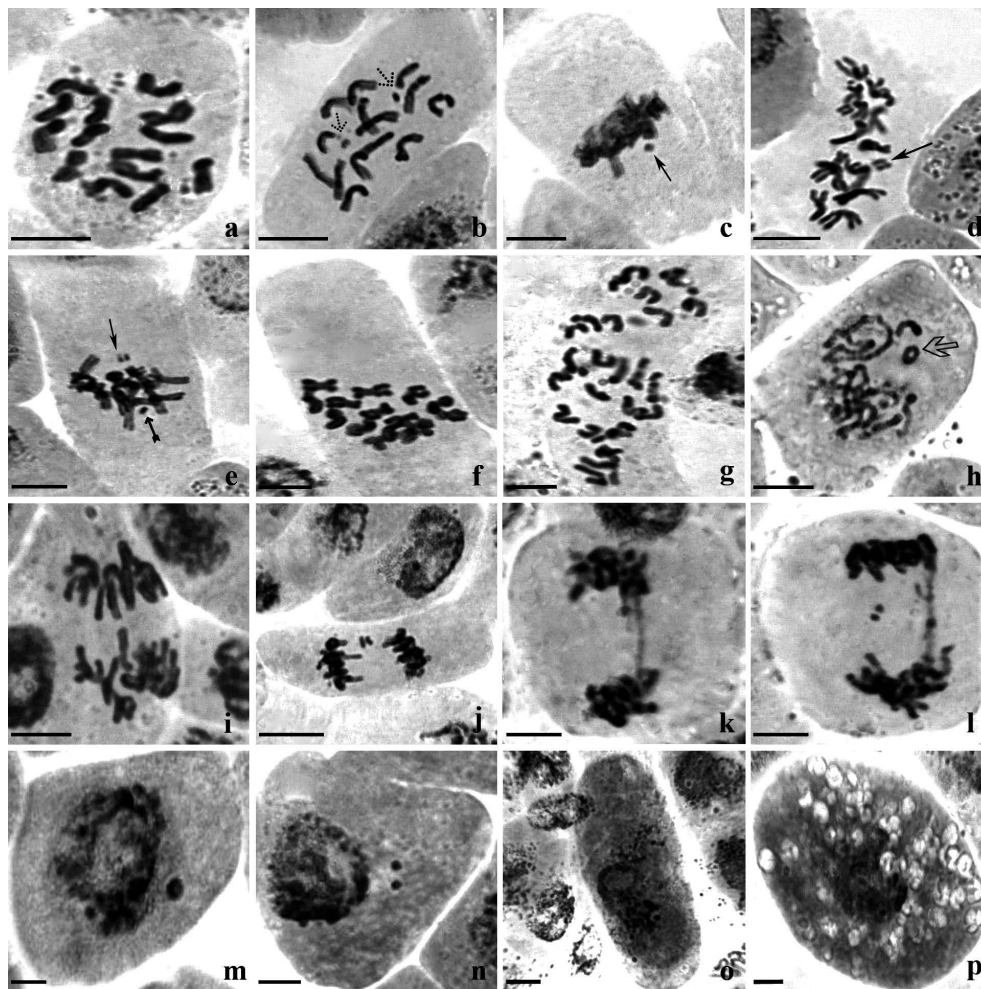


Fig. 1. NPs induced mitotic configurations at prophase (h), metaphase (a–g), anaphase (i–l) and resting stages (m–p). (a) $2n=14$. (b) Gaps (\leftrightarrow). (c) Stickiness with paired fragments (\leftrightarrow). (d) Paired fragments (\leftrightarrow). (e) Paired (\rightarrow) and unpaired (\rightarrow) fragments. (f) Pseudo-chiasmata. (g) Polyploidy $2n=28$. (h) Ring (\Rightarrow). (i) Multipolarity. (j) Paired fragments at anaphase. (k) Bridge. (l) Bridge with paired fragments. (m–n) Micronuclei. (o) Giant cell. (p) Vacuolated giant cell. Scale bar = $10\ \mu\text{m}$.

ered blockage of cells into mitosis as the most important cellular event after mutagenic treatment.

Cells with $2n=14$ (Fig. 1a) chromosomes are mostly observed in treatments. Stickiness of chromosomes and occasional laggard formation are studied in controls (Table 2). Types of aberration (Fig. 1b–p) observed in NP treatments are categorized in Table 2. They are possibly the outcome of breakages and interchanges, spindle malfunctioning and cellular metabolic disorders. Cells with paired fragments are notably observed in both metaphase and anaphase. Ghasem *et al.* (2011) performed karyotype analysis of different germplasms of *L. sativus* and reported mostly one and rarely two pairs of chromosomes with satellites. In the present study, paired fragments (Fig. 1c–e, j, l) may have arisen due to breakage of a satellite pair of chromosomes or breaks in identical site of chromatids. Ionizing radiations are reported to induce localized breakages in chromosomes possessing sub-terminal constrictions in *Nigella damascena* (Moutschen 1968) and *N. sativa* (Datta and Biswas 1983).

Micronuclei (one to two) of variable (Fig. 1m, n) sizes (condensed and uncondensed) are found to occur in NP treatments, suggesting their effective potentiality to induce chromosomal breakages. Giant cells with and without vacuoles (Fig. 1o, p) are observed in resting cells. Vacuolated giant cells are mostly noted in higher doses of NPs. Extensive vacuolation has been attributed to the cellular attempts of compartmentalizing NPs to reduce the phytotoxicity associated to its exposure (Li *et al.* 2014, Thakur *et al.* 2014). Both NPs induce similar type of responses with respect to mitotic aberrations.

Conclusion

The types of aberration observed in NP treatments are rather similar to the genotoxic effects induced following X-ray and gamma irradiations, suggesting the mutagenic potentiality of the studied nanoparticles. Genotype sensitivity of NP treatments is significant for potential effectiveness in inducing genetic variations.

Acknowledgements

The authors are thankful to Horticultural Research Station, Nadia, West Bengal for generous supply of the germplasm.

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