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VARIETAL DIFFERENCE IN NATURAL COLONIZATION BY ARBUSCULAR MYCORRHIZAL FUNGI AND ITS INFLUENCE ON PLANT CHARACTERS AND PHOSPHORUS NUTRITION IN SESAME

V.S. HARIKUMAR¹

Abstract. Twenty accessions of sesame (*Sesamum indicum* L.) plants were evaluated for arbuscular mycorrhizal (AM) colonization and its relationship with growth, yield and tissue P content. Sesame accessions exhibited variations in both frequency (%F) and intensity (%M) of colonization by AM fungi. Among the accessions, 30% had a comparatively higher %F whereas in other ones the %F was more or less on par. The %M was comparatively more in 40% of the accessions while two accessions had a very low %M. AM fungi which occurred in the rhizosphere of sesame accessions belonged to two species of *Acaulospora* and three species of *Glomus*. *Glomus* spp was found dominant in the rhizosphere soils of sesame. Correlation studies revealed a significant positive relationship between fungal variables as well as between fungal variables and plant characters such as growth, yield and tissue P content.

Keywords: AMF, host variety, plant characters, sesame.

INTRODUCTION

The ubiquitous arbuscular mycorrhizal (AM) fungi are an integral component of any soil system where they form obligate symbiosis with the roots of over 80% terrestrial plant species (van der Heijden & Sanders, 2002). Genetic variation within plant species can influence both the degree of root colonization by mycorrhizal fungi and the response of the plant to mycorrhizal symbiosis (Peterson & Bradbury, 1995). For example, Harikumar & Potty (2002) screened 257 genetic stocks of field grown sweet potato and found that 20% of them responded to natural colonization to the tune of 25% whereas, another 20% had 50% colonization in their root system and the remaining 60% had a very high level of colonization. Selection of host plant germplasm to maximize productivity and pest resistance or responsiveness to fertilization can affect the ability of the host to sustain or benefit from mycorrhizas (Kaeppeler *et al.*, 2000; Tawaraya, 2003).

Sesame (*Sesamum indicum* L., Fam. Pedaliaceae) is cultivated in tropical, subtropical and southern temperate regions of the world for its seeds which are a

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rich source of edible oil. Incidence of AM colonization in sesame roots and infective propagules in the rhizosphere of the crop have been reported earlier (Sulochana *et al.*, 2000; Harikumar, 2015). The evaluation of plant varieties for AM colonization and endophyte specificity helps to identify the most compatible symbiont with the host (Dhillon, 1992). The present paper reports the varietal differences and species specificity in AM colonization in 20 sesame varieties and its influence on plant characters.

MATERIALS AND METHODS

Twenty accessions of sesame (courtesy KAU (RS) Kayamkulam) collected from different parts of the country were evaluated for AM colonization and its relationship with growth, yield and P content of the plants. The study was conducted in a farmer's field located at Alappuzha, Kerala. The soil was sampled for analysis of physico-chemical properties (Jackson, 1973) and AM fungal population prior to the initiation of the experiment. The soil was a sandy Entisol with a pH 5.5 (1: 2:5; soil: water) and organic carbon 1.17%. Soil nutrient determinations included 108 kg N h⁻¹, 25 kg P h⁻¹ and 19.40 kg K h⁻¹. The total indigenous AM fungal spore density prior to the start of the study was 103 spores per 50 ml soil.

Seeds were hand sown in rows at a distance of 1 ft with three replicate pits for each accession. After emergence of seedlings, the plant number in each pit was thinned to three to avoid overcrowding. Plants did not receive any fertilization or irrigation throughout the growth period. Weeds were controlled by hand as required.

Plants were harvested at 75 days after emergence. Three replicated plants for each accession were dug out with almost the entire root system intact. Rhizosphere soil samples were collected from 10–20 cm depth. The harvested plants were utilized for monitoring mycorrhizal colonization, growth and yield. The tissue P content was determined as per Jackson (1973).

The undamaged fine roots of test plants were cut into 1 cm root segments from these about 50 segments were selected at random. The root segments were thoroughly washed and stained with cotton blue (Phillips & Hayman, 1970). The root segments were mounted on clean microscopic slides in a mixture of glycerol and lactic acid (v/v). The root segments were gently squashed and covered by a glass cover slip and observed under a compound microscope (Nikon Eclipse E 400) using different magnifications. Mycorrhizal variables, frequency (%F) and intensity (%M) of AM colonization were calculated following the method of Trouvelot *et al.*, 1986). Spores were extracted from a sub-sample (50 ml) of each rhizosphere soil sample by wet-sieving and sucrose density gradient centrifugation (Daniels and Skipper, 1982) and examined using a Zeiss Stemi-DV4 stereomicroscope. Spores of each morphotype were mounted on slides in polyvinyl alcohol-lactic acid-glycerol (PVLG) (Koske & Tessier, 1983) and PVLG mixed 1:1 (v/v) with Melzer's reagent. Spores were examined using a Nikon Eclipse 400 research microscope and identified up to species level using the manual for the identification of VA mycorrhizal fungi by

Schenck & Pérez (1990). Spore density was determined as the number of healthy appearing spores per 50 ml soil. Isolation frequency was calculated as the percentage of samples in which the particular genus or species was present.

RESULTS AND DISCUSSION

AM colonization in sesame accessions exhibited variation in both %F and %M. Among the 20 accessions screened, 30% had a comparatively higher (>40%) %F whereas in other ones the %F was more or less on par. %M was comparatively more (>8%) in 40% of the accessions while two accessions (SI 68, SI 69) had a very low %M (Table 1). While AM fungi are known to vary in their ability to colonize and transfer P to the plant and confer other beneficial effects, little is known to the exact role of the host genotype in the expression of AM fungi. In the present study, though the accessions were grown in the same field under identical environmental condition, there was significant variation in root colonization by AM fungi which could be ascribable at least in part to genetic and physiological factors controlling host/fungus compatibility. Recently discovered mycorrhizal mutants (myc⁻) of pea offer great promise for the study in this direction (Graham & Eissenstat, 1994). Similar genotype dependent variation for AM fungal colonization has been reported in wheat (Azcón & Ocampo, 1980) sorghum (Clark, 1983) pearl millet (Krishna *et al.*, 1985) coconut (Thomas & Ghai, 1987) and cowpea (Mercy *et al.*, 1990).

AM fungi which occurred in the rhizosphere of sesame accessions belonged to two species of *Acaulospora* and three species of *Glomus* (Table 1). *Glomus* spp was found dominant in the rhizosphere soils. The results revealed that *G. mosseae* associated with most accessions with the highest frequency (Figure 1). *Glomus* species are capable of colonizing roots *via* fragments of mycelium or mycorrhizal root pieces (Daniell *et al.*, 2001). This may be the possible reason for the predominance of the genera over other genera in most agricultural soils (Morton, 1988). Furthermore, they are widely adaptable to the varied soil conditions and survive in acidic as well as alkaline soils (Pande & Tarafdar, 2004).

Correlation studies between fungal variables (%F, %M) revealed that there exists a significant positive relationship between these variables. A similar significant positive correlation was observed between fungal variables and plant characters (Table 2). %F was significantly correlated with measured variables such as growth, yield and P content. %M also followed the same pattern of relationship. Harikumar & Potty (2002) in a field study with 257 genetic stocks of sweet potato for mycorrhizal colonization and its response on growth of the crop observed that fungal characters showed a positive relationship with underground plant characters (root and tuber) only while the above ground portion had no direct bearing on the interaction. By contrast, in the present study all the parameters examined showed a significant correlation with the fungal variables further confirming the importance of indigenous AM endophytes on the growth and nutrition of sesame.

Table 1

Frequency (%F) and intensity (%M) of root colonization and AMF species associated with sesame accessions

Accession no.	Colonization*		AM Species**
	%F	%M	
SI 2	43.33 ^{bcd}	8.70 ^c	3, 4, 5
SI 7	43.33 ^{bcd}	9.17	3, 4, 5
SI 15	26.67 ^{ef}	5.66 ^c	2, 4
SI 17	23.33 ^{ef}	4.50 ^{ef}	2, 4
SI 25	26.67 ^{ef}	5.90 ^{de}	2, 3
SI 30	56.67 ^{ab}	17.70 ^a	1, 3, 4, 5
SI 32	46.67 ^{bc}	8.17 ^{cd}	1, 4, 5
SI 37	28.33 ^{def}	4.00 ^{ef}	2, 4
SI 42	28.33 ^{def}	4.93 ^{ef}	3, 5
SI 47	68.33 ^a	19.30 ^a	1, 3, 4, 5
SI 48	50.00 ^b	14.50 ^b	1, 3, 4
SI 56	33.33 ^{cde}	8.90 ^c	4, 5
SI 57	33.33 ^{cde}	8.60 ^c	2, 4
SI 58	26.67 ^{ef}	5.43 ^{ef}	3, 5
SI 59	26.00 ^{ef}	5.27 ^{ef}	3, 4
SI 63	23.33 ^{ef}	4.50 ^{ef}	2, 3
SI 66	23.33 ^{ef}	5.07 ^{ef}	2, 3
SI 68	16.67 ^f	3.07 ^f	3
SI 69	28.33 ^{def}	3.63 ^{ef}	4, 5
SI 70	28.33 ^{def}	4.13 ^{ef}	4, 5

* Means in each column with different letters are significantly different ($P < 0.05$) by Tukey's HSD

** 1. *A. delicata* 2. *A. lacunosa* 3. *G. dimorphicum* 4. *G. mosseae* 5. *G. versiformae*

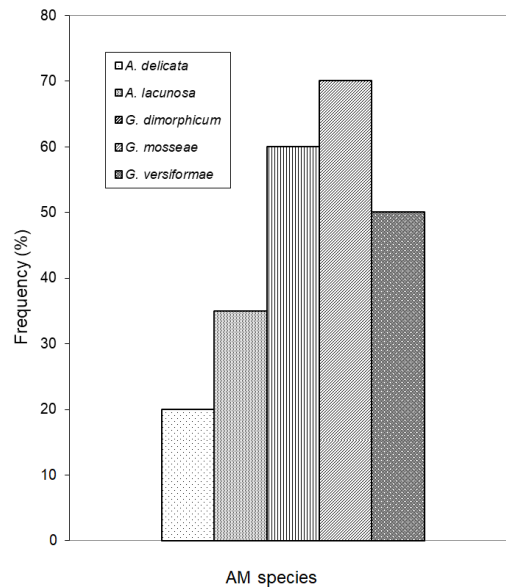


Fig. 1. Frequency of AM species in the rhizosphere of sesame accessions.

Table 2

Frequency (%F) and intensity (%M) of root colonization and AMF species associated with sesame accessions

Parameters	Mycorrhizal parameters	
	% F	% M
Mycorrhizal parameters		
% F	1.000	
% M	0.897***	1.000
Growth characters		
Root length (cm)	0.479***	0.547***
Shoot length (cm)	0.499***	0.528***
Leaf Number plant ⁻¹	0.707***	0.781***
Biomass production		
Root fresh weight (g)	0.593***	0.669***
Shoot fresh weight (g)	0.493***	0.563***
Total biomass (g)	0.538***	0.614***
Dry matter production		
Root dry weight (g)	0.573***	0.667***
Shoot dry weight (g)	0.526***	0.621***
Total dry weight (g)	0.544***	0.641***
Yield		
Pod Number (plant ⁻¹)	0.404***	0.358***
P efficiency		
P content (mg g ⁻¹)	0.546***	0.480***

*** $P < 0.001$.

CONCLUSION

Sesame accessions revealed AM colonization attributable to the species of *Acaulospora* and *Glomus* which varied with varieties. In general, both %F and %M were low in sesame accessions. Nevertheless, the crop is greatly dependent on indigenous AM fungi for growth and nutrition as a significant positive relationship between plant and fungal characters could be drawn out through correlation analysis.

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RED LISTED LICHEN SPECIES WITHIN OLD GROWTH AND YOUNG GROWTH FORESTS FROM ROMANIA

I. VICOL¹

Abstract. In the studied areas of a total of 19 red listed lichen species from Romania there were observed 4 red listed lichen species, for instant: *Cetraria islandica* (L.) Ach. (1803), *Cetraria saepincola* (Ehrh.) Ach. (1803), *Hypotrachyna sinuosa* (Sm.) Hale (1975), and *Lobaria pulmonaria* (L.) Hoffm. (1796). The red listed lichen species were predominantly identified in mountainous areas within old growth forests well conserved, the most of them being protected areas. As regards young growth forests, only one red listed lichen species was found in a mixed forest represented in a great deal by ancient oaks from the Romanian Plain. In the field it was observed that red listed lichen species have been found especially on old trees, such as: oak and beech. In the lowland areas, within mixed forests represented especially by oaks *Hypotrachyna sinuosa* has been identified while within montane forests well represented by beech, fir and spruce the presence of *L. pulmonaria* has been observed. In addition to forest habitats, *C. islandica* and *C. saepincola* were found on mugo pine and common juniper in subalpine and alpine belts with undetectable anthropogenic pressure.

Keywords: forest, lichen, mountainous areas, oak, beech.

INTRODUCTION

At European level both old growth forests and their characteristic lichen species are nowadays relictual (Scheidegger & Goward, 2002). The mixed forests from Europe are predominantly represented by oak, elm and beech which have been harvested and converted into secondary forests. The vast forestlands are threatened because of increasing of agricultural, urbanization and industrial activities; in this regard to avoid the loss of the forest biodiversity protected areas have been designed (Nascimbene *et al.*, 2013).

It has been estimated that 0.2% of mixed forests from Europe are old-growth, the other being intensively managed for economical development. Nevertheless, Romania is very well represented by deciduous forests compared with western European countries (Nascimbene *et al.*, 2013).

Old oaks host a rich lichen species diversity due to their favourable microhabitats, but in young growth forests the lichen diversity is subjected to a

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progressive loss depending on the scale of the shade degree (Paltto *et al.*, 2011). In Europe, during forest management, the old oaks were retained constituting a network of the individual patches whose elements act an important role both to dispersal diaspores and to lead further a spatial model for conserving lichen species diversity including many red listed species (Ranius *et al.*, 2008). Within forest habitats from Swedish, on deciduous trees older than 150 years were found lichen species associated to old-growth forests, for instance: *Calicium salicinum*, *Lobaria pulmonaria* and *Mycobilimbia sabuletorum*. These lichen species are considered “late-successional species” probably favoured by texture and chemical composition of the tree bark. Forest continuity includes besides old trees other old growth forest elements such as “number of tree layers” and “stand age”; therefore to conserve the old forest attributes is necessary to retain groups of old trees during clear-cutting (Gustafsson *et al.*, 1992). Although a great attention must be paid to the anthropogenic impact that is responsible to decreasing the red listed lichen species in protected areas, considered nowadays as major refuge habitats (Motiejūnaitė, 2011).

Within young-growth forests there are predominant common lichen species which belong to *Xanthorion* group unlike to sensitive lichen species found in old-growth forests. The sensitive lichen species are affected by forest management including clearing and cutting, the density of young trees and the presence of shrubs; therefore “higher habitat quality requirements” as characteristic environment for them are necessary (Aragón *et al.*, 2010).

The aim of this study consists in highlighting the importance of the presence of red listed lichen species within young-growth and old-growth forests. The objectives of the study are the following: (1) finding out the presence of red listed lichen species within young-growth and old-growth forests and (2) revealing the presence of red listed lichen species within forest habitats situated in geomorphological units such as: plain, hills and mountains.

MATERIALS AND METHODS

The field researches were performed during 2009–2015 within old growth and young growth forests from Alba, Bihor, Botoşani, Braşov, Caraş-Severin, Cluj, Gorj, Hunedoara, Ilfov, Neamţ, Prahova, Sibiu and Vaslui counties (Fig. 1).

The specimens collected from the field were determined in laboratory using keys (Moruzi & Toma, 1971; Purvis *et al.*, 1994; Ciurchea, 2004), stereomicroscope (Zeiss Stereo CL 1500 ECO), optical microscope (Zeiss Scope A1) and chemical reagents such as IIK (iodine-potassium iodide) and KOH (potassium hydroxide).

Identification and nomenclature of the tree species is according to www.uk.ipni.org, whilst for lichen nomenclature it was used www.mycobank.org.

The identified material is preserved within the Mycological Herbarium, Lichen Collection (BUCML) Institute of Biology, Romanian Academy, Bucharest, Romania.

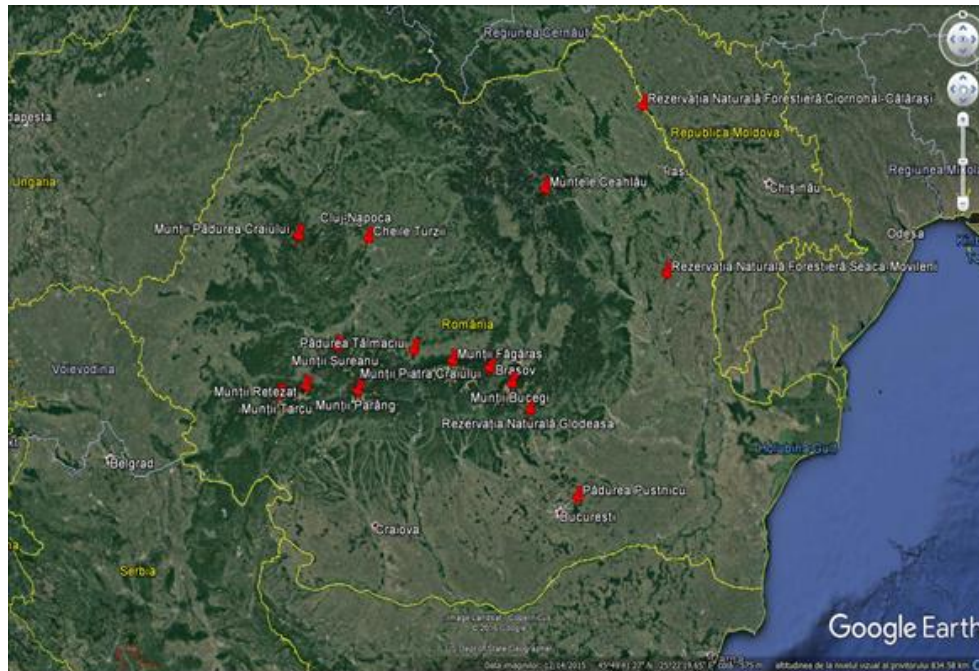


Fig. 1. The studied sites within red listed lichen species were identified.

RESULTS

Within the investigated forests there were identified 4 lichen species which are found on the Romanian Red List of lichens (Sârbu *et al.*, 2007; Ardelean *et al.*, 2013), for instance: *Cetraria islandica* (L.) Ach. (1803), *Cetraria saepincola* (Ehrh.) Ach. (1803), *Hypotrachyna sinuosa* (Sm.) Hale (1975), and *Lobaria pulmonaria* (L.) Hoffm. (1796).

The chorology of the identified red listed lichen species is the following:

Cetraria islandica (L.) Ach. (1803) Fig. 2.

Alba County: Șureanu Mountain, at the end of the ski lifts, found on terricolous substrata, in a meadow, leg. Vicol Ioana, 24.10.2012, det. Vicol Ioana, 31.10.2012, BUCM L1749; found within shrubs represented by *Juniperus communis* Thunb. (1784), leg. Vicol Ioana, 24.10.2012, det. Vicol Ioana, 08.11.2012, BUCM L1758.

Brașov County: Pietra Craiului Mountains, Turnu Peak, found on terricolous substrata, leg. Vicol Ioana, 27.06.2012, det. Vicol Ioana, 03.07.2012, BUCM L1748.

Caraș-Severin County: Țarcu Mountains, Țarcu Peak, northern exhibition, alt. 1900 m, found on saxicolous substrata, within shrubs represented by *Vaccinium* sp., leg. Ion Roxana Georgiana, 06.08.2013, det. Vicol Ioana, 09.05.2014, BUCM L2292.

Gorj County: Parâng Mountains, near Rânca locality, found on terricolous substrata, in a meadow, leg. Vicol Ioana, 23.10.2012, det. Vicol Ioana, 02.11.2012,

BUCM L1750, BUCM L1753, BUCM L1754, BUCM L1755, BUCM L1757; Parâng Mountains, between Râncă and Obârșia Lotrului, found on terricolous substrata, in a meadow, leg. Vicol Ioana, 23.10.2012, det. Vicol Ioana, 01.11.2012, BUCM L1751, BUCM L1752; leg. Vicol Ioana, 23.10.2012, det. Vicol Ioana, 31.10.2012, BUCM L1756.

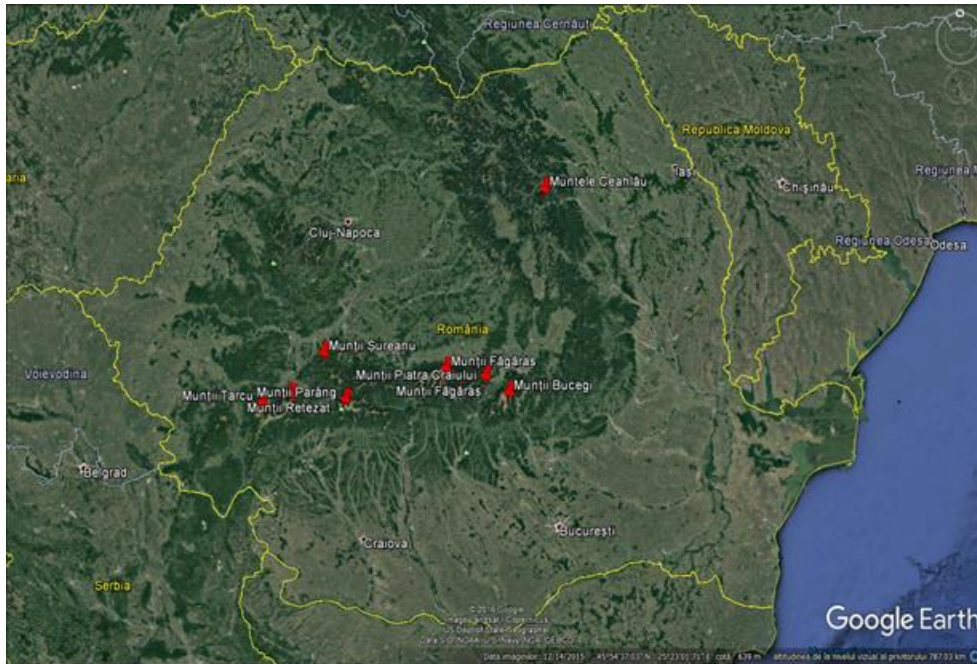


Fig. 2. The chorology of the red listed *Cetraria islandica* in Romania.

Hunedoara County: Retezat Mountains, Retezat National Park, found on saxicolous substrata, leg. Vicol Ioana, 29.05.2013, det. Vicol Ioana, 10.06.2013, BUCM L1946; Culmea Lolaia, found on terricolous substrata, leg. Vicol Ioana, 29.05.2013, det. Vicol Ioana, 03.06.2013, BUCM L1963; Râușor, Colții Prelucelor, found on terricolous substrata, leg. Ion Roxana Georgiana, det. Vicol Ioana, 23.09.2013, BUCM L2009; Culmea Prelucelor, found on terricolous substrata, leg. Vicol Ioana, 21.08.2014, det. Vicol Ioana, 26.08.2014, BUCM L2375; Ștevia Valley, eastern exhibition, alt. 1847 m, found on terricolous substrata, leg. Vicol Ioana, 20.08.2014, det. Vicol Ioana, 26.08.2014, BUCM L2488; towards Ștevia Valley, in a beech and spruce forest, lat. 45.39756°N, long. 22.84426°E, alt. 1775 m, found on saxicolous substrata, leg. Vicol Ioana, 20.08.2014, det. Vicol Ioana, 15.09.2014, BUCM L2499.

Neamț County: Ceahlău National Park, Ceahlău Mountain, near Toaca Peak, found on terricolous substrata, in subalpine plateau within juniper shrubs, leg. Vicol Ioana, 13.10.2011, det. Vicol Ioana, 18.10.2011, BUCM L1561, BUCM

L1562; near Toaca Peak, found on terricolous substrata, leg. Vicol Ioana, 25.07.2014, det. Vicol Ioana, 08.08.2014, BUCM L2380; alpine plateau, found on terricolous substrata, leg. Vicol Ioana, 26.06.2014, det. Vicol Ioana, 26.06.2014, BUCM L2351; tourist route Durău - Duruitoarea Waterfall - Dochia Chalet, Polița cu ariniș, alt. 1624 m, found on terricolous substrata, leg. 26.06.2014, BUCM L2365; La Pălărie, found on terricolous substrata, leg. Vicol Ioana, 23.07.2014, det. Vicol Ioana, 07.08.2014, BUCM L2425.

Prahova County: Bucegi Mountains, found on terricolous substrata, leg. Mogâldea Daniela, 11.07.2011, det. Vicol Ioana, 26.09.2011, BUCM L1560.

Sibiu County: Făgăraș Mountains, Bâlea Lake, alt. 2044 m, lat. 45°36'11.13''N, long. 24°36'45.95''E, found on saxicolous substrata, in an alpine grassland, leg. Fiera Cristina, 05.09.2015, det. Vicol Ioana, 24.09.2015, BUCM L2652.

Cetraria saepincola (Ehrh.) Ach. (1803) Fig. 3.

Hunedoara County: Retezat Mountains, Retezat National Park, Culmea Prelucele, alt. 1849, found on *Pinus mugo* Turra (1764), leg. Vicol Ioana, 21.08.2014, det. Vicol Ioana, 07.08.2014, BUCM L2492.

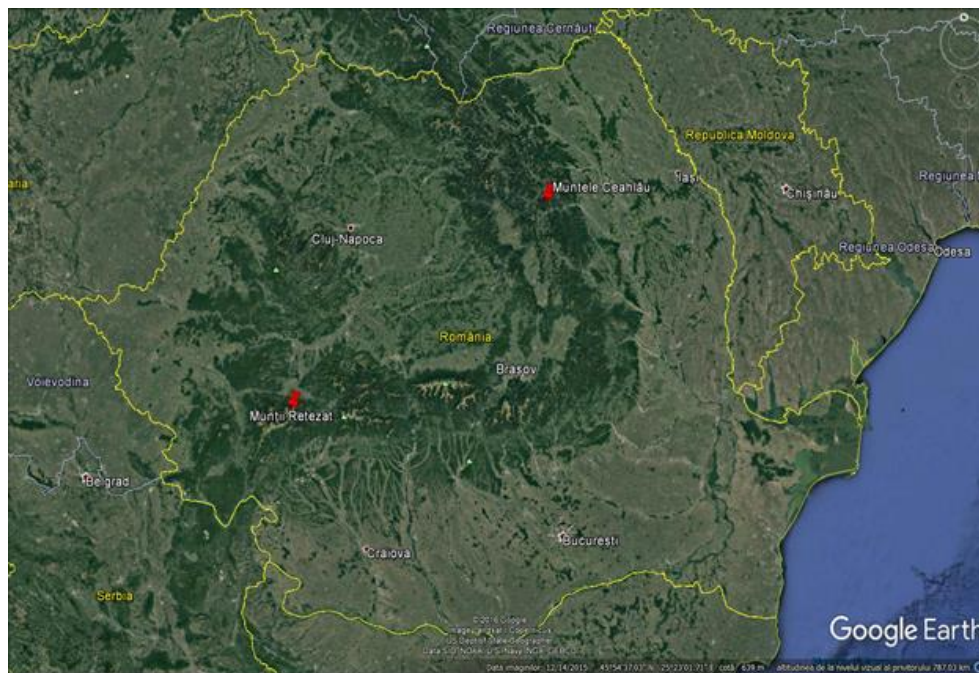


Fig. 3. The chorology of the red listed *Cetraria saepincola* in Romania.

Neamț County: Ceahlău Mountain, Ceahlău National Park, La Pălărie, alt. 1743 m, on *Juniperus communis* Thunb. (1784), leg. Vicol Ioana, 23.07.2014, det. Vicol Ioana, 07.08.2014, BUCM L2449 (Vicol, 2015b).

Hypotrachyna sinuosa (Sm.) Hale (1975) Fig. 4.

Botoșani County: Ciornohal-Călărăși Nature Reserve Forest, found on *Quercus robur* L. (1753), leg. Vicol Ioana, 16.07.2013, det. Vicol Ioana, 17.09.2013, BUCM L2081 (Vicol, 2016).

Cluj County: along a trail between Turzii Gorges Chalet towards Turzii Gorges, found on *Quercus* sp., leg. Vicol Ioan, 18.04.2013, det. Vicol Ioana, 13.05.2013, BUCM L1930.

Ilfov County: Pustnicul Forest, found on *Quercus cerris* L. (1753), leg. Vicol Ioan, 27.03.2009, det. Vicol Ioana, 22.06.2009, BUCM L1363; leg. Vicol Ioan, 30.04.2009, det. 23.06.2009, BUCM L1389 (Vicol, 2010).

Sibiu County: Tâlmaciu Forest, on saxicolous substrata, leg. Vicol Ioana, 28.03.2011, det. Vicol Ioana, 04.04.2011, BUCM L1616.

Vaslui County: Seaca-Movileni Nature Reserve Forest, found on *Quercus pedunculiflora* K. Koch., leg. Vicol Ioan, 17.05.2012, det. Vicol Ioana, 30.05.2012, BUCM L1810 (Vicol, 2015a).

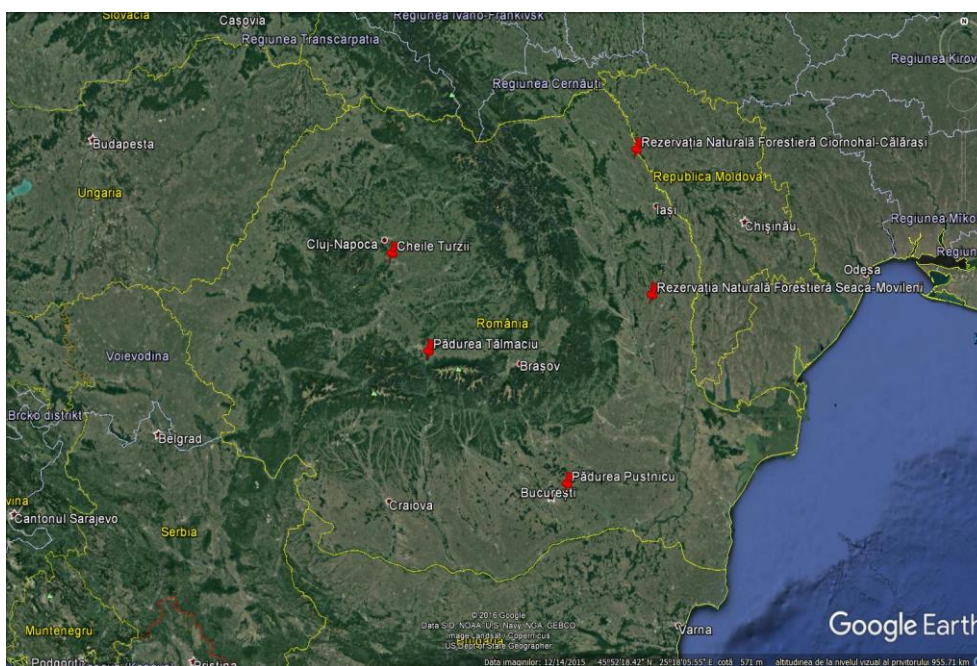


Fig. 4. The chorology of the red listed *Hypotrachyna sinuosa* in Romania.

Lobaria pulmonaria (L.) Hoffm. (1796)

Bihar County: Apuseni Natural Park, Pădurea Craiului Mountains, between Izvorul Minunilor (Stâna de Vale) and Culmea Muncelu, found on *Fagus sylvatica* L. (1753), leg. Vicol Ioana, 30.08.2014, det. Vicol Ioana, 02.09.2014 BUCM L2477.

Neamț County: Ceahlău Mountain, Ceahlău National Park, near Izvorul Muntelui locality, found on *Fagus sylvatica* L., 13.08.2014, alt. 817 m, lat. 46.95374°N, long. 25.98634°E (Vicol, 2015b).

Prahova County: Glodeasa Nature Reserve, found on corticolous substrata, leg. Ion Constanța Mihaela, 13.05.2009, det. Vicol Ioana, 16.07.2009, BUCM L1414; Bucegi Mountains, tourist route Căminul alpin – Munticel – Poiana Coștilei – Pichetul Roșu – Cabana Mălăiești, alt. 1288 m, lat. 45.43060°N, long. 25.50908°E, found on *Fagus sylvatica* L. (1753), leg. Vicol Ioana, 04.06.2014, det. Vicol Ioana, 19.06.2014, BUCM L2312; Bușteni, Poteca Munticelului, found on *Fagus sylvatica* L., 04.06.2014, alt. 1288 m, lat. 45.43060°N, long. 25.50908°E, exhibition N-V.

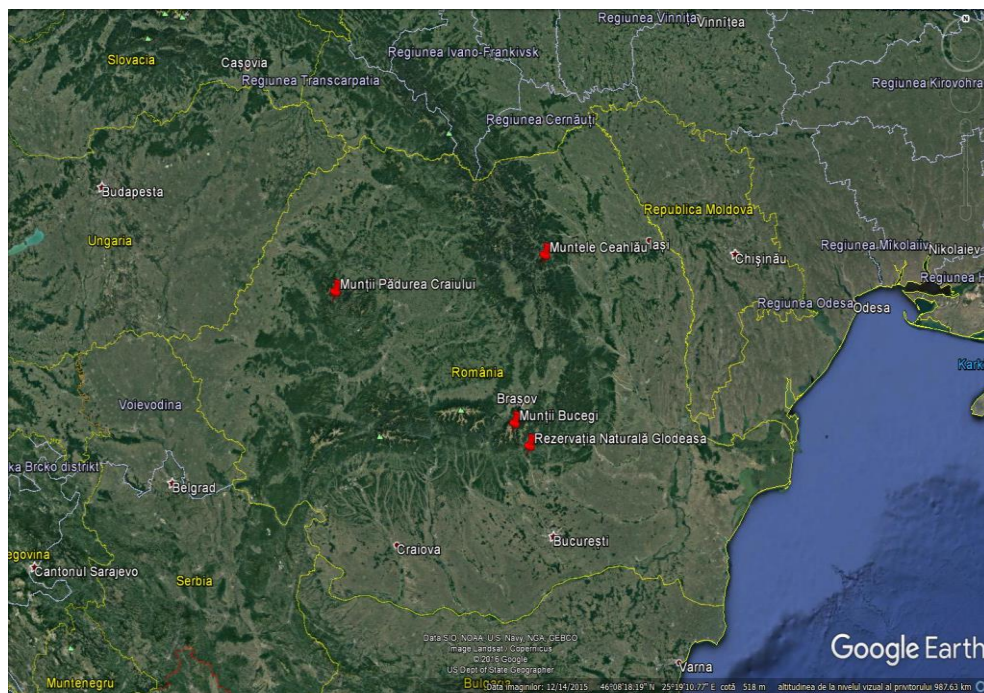


Fig. 5. The chorology of the red listed *Lobaria pulmonaria* in Romania.

Based on the above chorology, the red listed lichen species were identified in both plain, hilly and mountainous areas. Thus, the red listed lichen species were found to a great deal in mountainous areas represented by old growth forests (Table 1). These forests are predominantly represented by beech accompanied by fir and spruce.

Table 1

The presence of red listed lichen species in old growth forests

plain area	hilly area	mountainous area
–	<i>Hypotrachyna sinuosa</i>	<i>Cetraria islandica</i>
–	–	<i>Cetraria saepincola</i>
–	–	<i>Lobaria pulmonaria</i>

In the case of young growth forests situated in plain areas from the Romanian Plain, near the eastern part of the Bucharest Municipality only one of the red listed lichen species has been found, namely *H. sinuosa*. Regarding the young growth forests from hilly areas none of the lichen species has been found (Table 2). The young growth forests are mainly characterised by the predominance of the oak.

Table 2

The presence of red listed lichen species in young growth forests

plain area	hilly area	mountainous area
<i>Hypotrachyna sinuosa</i>	–	–
–	–	–
–	–	–

At the upper limit of forest, in subalpine and alpine areas where anthropogenic impact is low, within meadows and shrubs represented by mugo pine and common juniper were found out *C. islandica* and *C. saepincola*. Also, these species were found within old coniferous forests from mountainous areas.

DISCUSSION

As regards the research theme there are least concerns on red listed lichen species and their associated forest habitats in Romania. A larger number of the red listed lichen species have been found in mountainous areas due to a lowest anthropogenic impact. In both inferior, subalpine and alpine belts especially within protected areas the number of red listed lichen species is greater due to conservation of their specific microhabitats. These microhabitats are represented among other characters by large circumferences of the tested trees. Thus, in Neamţ County, Ceahlău National Park, one of the red listed lichen species (*L. pulmonaria*) has been identified on *F. sylvatica* with a circumference of 3.20 m. This is one of the most relevant particularities of old growth forests because they highlight the ancient forest continuity (Nascimbene *et al.*, 2007). *Lobaria pulmonaria* is related to old growth forests especially mountainous beech forests where this species is more frequently due to suitable environmental conditions (Svoboda *et al.*, 2011). The absence of cyanolichens from young-growth forests is caused by intensive management of forestry practices (Nascimbene *et al.*, 2007). In other European countries, *L. pulmonaria* is found in Red List of Estonia (Randlane *et al.*, 2008).

According to field observations, Pustnicul Forest is a matrix with old trees and planted trees. The old ones were integrant elements of the old growth forests; therefore, one of the red listed lichen species has been found on an old oak (a circumference of 0.91 m), namely, *H. sinuosa* (Vicol, 2010). In the case of this species the vegetation layers of forest act as biotic barrier against atmospheric pollution. The Pustnicul Forest is situated in the eastern part of Bucharest, intensively polluted by industrial and car traffic activities (Vicol, 2014). The lichen species which are rare have a long lifetime and a deficient dispersal; therefore these species depend on their specific habitats as are old growth forests with a low intensity of the human impact (Scheidegger & Goward, 2002). As regards the trees pattern from Pustnicul Forest, which is an intensively managed forest, due to its ecological continuity the red listed lichen species are favoured. Otherwise, the presence of the one red listed lichen species within this forest is mainly caused by its fragmentation (Svoboda *et al.*, 2011). The ecological performance of red listed lichen species in a young growth forest mixed with relictual oaks consists in a selective cutting of trees, so that to create a matrix with young and old trees, in which the dispersal of the propagules to be efficient (Nascimbene *et al.*, 2007; Morley & Gibson, 2010). The lichen species are affected by attributes of old growth forests, such as circumferences of host trees, tree species diversity (Mežaka *et al.*, 2008; Vicol, 2016) and rhytidome crevice depth (Vicol, 2015b).

It is important to retain that *Lobaria pulmonaria* is a “flagship species” and a “signal species for rapidly assessing the conservation importance of sites” (Nascimbene *et al.*, 2007) whilst oaks are “keystone” (Ranius *et al.*, 2008) with a great significance for native forests conservation in Romania (Negrean & Ciortan, 2014).

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LICHENS – SOME RELEVANT ASPECTS OF THE *IN VITRO* CULTURE AND ULTRASTRUCTURAL PECULIARITIES

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Abstract. This paper summarises “*in vitro*” reactivity and evolution of *Xanthoria parietina* Beltr. and *Cetraria islandica* explants, biotechnological potential assessment of in situ and *in vitro* symbiotrophic formations of *Cetraria islandica* (L), the main aspects of lichen symbionts ultrastructure for the *Xanthoria parietina* Beltr., *Pseudoevermia furfuracea* species, and *Usnea barbata* L. Mott.

The investigations presented in this paper focused mainly on the possibility to elaborate alternative efficient methodologies for lichen species culture and artificial biomass resynthesis (symbiotrophic formations), using “*in vitro*” systems.

Keywords: mycobiont, ascospore isolation, lichen culture, symbiotrophic formations, resynthesis, ascospore isolation.

INTRODUCTION

As we know from the literature, lichens are the expression of the union between a photosynthetic partner and a heterotrophic one, the photobiont and mycobiont, respectively.

Their close interaction creates a new organism with distinct features that allows them to survive in inappropriate environments for each other (Zarnea & Popescu, 2011). As a result of symbiotic relation, the structure of the partners undergoes a series of changes (Moya *et al.* 2015).

The controversial nature of lichens stimulated the interest of many researchers for their study, on multiple levels. In this context, a history and classification of the experimental approaches in lichenology is exposed by Lobakova S. and Smirnov A. (2012).

Many bioactive compounds are unique for lichens and exert an antimicrobial and/or antitumoral (usnic acid, for example) action (Grujicic *et al.* 2014.) On the other hand, a considerable number of species became endemic, a good reason for their bioconservation and repopulation strategies.

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At the same time the sensibility of some lichen species to the environmental changes recommends them as indicators of life medium quality (Vicol 2011 a).

The aim of our researches was the development of some alternative methodologies of bioconservation, resynthesis and biomass obtaining with a significant content of bioactive substances, using the advantages offered by “in vitro” systems and ultrastructural analysis of symbiosis partners.

MATERIALS AND METHODS

As biological material for study we used the following species: *Usnea barbata* (L.) Weber ex Wigg., *Cetraria islandica* (L.) Ach., *Xanthoria parietina* (L.) and *Pseudoevernia furfuracea* L.

Some characteristics of them like pharmacological value, the abundance in Romania lichenobiota besides the few data existing in the national scientific literature are arguments for our studies.

The protocols were adapted according to the literature (Ahmadjian, 1993), (Honegger, 1993), (Stocker – Worgotter, 2002), (Yamamoto *et al.*, 2002) and (Yoshimura *et al.*, 2002) to our objectives, considering the culture medium, the inoculum type and incubation conditions.

We tested a wide range of culture media (liquid and solid, organic and mineral, synthetic and naturals) water – agar 2% (WA2) (Yamamoto *et al.*, 1985); Murashige & Skoog (Ahmadjian, 1961), Malt – Yeast extract (MY) (Ahmadjian, 1961); nutritive organic medium, Trebouxia (Tonm) (Ahmadjian, 1967); Honegger (Honegger, 1993); Honegger with Gelrite; Knop; Knop with bark extract; Knop with soil extract, BBM, BBM – modified.

The grinding paste and derived from this, the “whole suspension” and micropipettes method was used for algal green isolation (Friedl *et al.*, 2008) and for resynthesis experiments.

For mycobiont isolation, two sources have been used, namely thallus explants, for *Cetraria islandica* and *Xanthoria parietina*, and spores from apothecia, dispersed directly on the surface of the MEYE (Ahmadjian, 1967) or MLB (Lilly & Barnett, 1951) for *Usnea barbata* and *Xanthoria parietina* species.

All parameters have been modified according to the pursued objectives.

Cryopreservation was a less used method for lichens. Banciu and Cristian, 2015, performed this method for *P. furfuracea*. The cryoprotection treatment used a solution of liquid half mineral concentration MS medium (Murashige & Skoog, 1962) supplemented with 6% (w/v) sucrose, and two variants of cryoprotectants: 5% Dimethyl sulfoxide (DMSO) and 5% glycerol, respectively 10% dimethyl sulfoxide (DMSO) and 10% glycerol, applied each for 30 minutes to 10 samples of lichens (Withers & Williams, 1985). A controlled cooling rate machine (from CryoLogic) was used for different cooling processes, with 2° C/min to 0° C, then

with 1° C/min to – 6° C, 0.3° C/min to – 32°, with 0.5° C/min to – 42° C, followed by immersion in liquid nitrogen (at – 196° C).

Also, we performed antioxidant activity (DPPH radical scavenging activity of extracts) and extraction of phenolic compounds for *Cetraria islandica*, after the method proposed by Marxen *et al.*, 2007.

The biotechnological potential using Folin – Ciocalteu reagent (Cristian D., Mitoi M., Brezeanu A., 2013), of *in vitro* cultures was exploited by determining flavonoids concentration.

Cultures evolution was monitored by macroscopic and microscopic analyses. The sections were examined in an TEM electron microscope – Philips – 200 (Toma N., Voicu D., Toma F.A., 2007) and EM – 125 (Selmi-Ucraina) electron microscope (Brezeanu A., Voicu D., 2008).

The symbiotrophic formations were cytologically analysed on squash samples by phase contrast as well as on semifine sections (after inclusion in synthetic resins – Epon 812) by photonic microscope (Scope A1, Zeiss model). For TEM electron microscopical analysis ultrathin technique which involves fixation in 3% glutaraldehyde and 1% OsO₄ at 4° C over night and inclusion in synthetic resin Epon 812 using the method proposed by Mascoro and Bozola, 2007, modified by us in proper formulas (Brezeanu, Voicu, 2008).

RESULTS AND DISCUSSION

Grinding paste methodology was used for all the following research directions.

The culture media which sustained the objectives were diverse.

Mainly the synthetic balanced culture media (BBM, the modified BBM , modified MS, Honegger) sustained artificial lichen thallus resynthesis experiments (Voicu D., Brezeanu A., 2008).

These satisfied nutritional requirements of both partners – the heterotroph acidophilic mycobiont and photoautotrophic and neutro – basophil photobiont; all three species investigated had a propitious development.

Guzow-Krzeminska & Stocker-Wörgötter E. (2013) also used modified BBM medium (enriched with mannitol), this enabling the balanced growth of both bionts.

The use of soil substratum was sustained, too, by the most of the lichenologists (Bubrick P., 1988b, Stocker – Wörgötter E., Wörgötter E. 2001), allowing us to conclude that a slightly enriched BBM medium could be used as a general medium for resynthesis experiments. The organic substances from the cortex stimulate algal cells proliferations; this fact is supported by the heterotrophic nature of *Trebouxia* in some cases (Ahmadjian, 1993).

For mycobiont lichens isolation the best source are the spore from apothecia released directly on the surface of the MEYE culture medium (Ahmadjian, 1967) for *Xanthoria parietina* or liquid BBM for *Usnea barbata* (L.) Mott. (Figs. 1, 2).



Fig. 1. The 6 days white column of the hyphae and 9 days mycobiont (Cristian D., Brezeanu A., 2013).

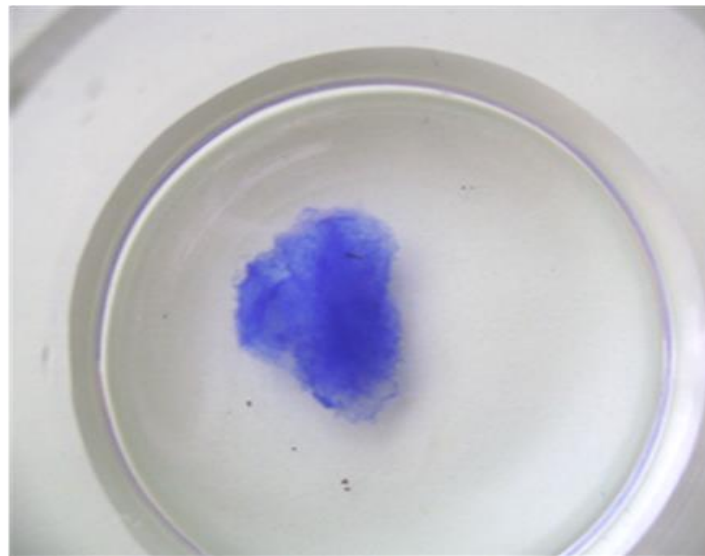


Fig. 2. *Usnea barbata* mycobiont culture in BBM-liquid, blue-cotton coloured (orig.).

The cultures evolution is stimulated by an alternative regime of light and dark with equal periods (12 hours) while continuous light has an inhibitory character.

Increasing the period of darkness stimulates the mycobiont development and thus decreases symbiotic interactions.

High temperatures (20–26°C) inhibit the symbiotic evolution of cultures while the low ones (12–15°C) stimulate symbiotic interaction between mycobiont and photobiont.

The explanation lies in the inhibitory partial character of low temperatures upon the nutrition capacity of mycobiont and photosynthetic yield that is forced to establish symbiotrophic relations without which it is doomed.

The periods necessary for symbiotrophic interactions differed according to species, photobionts density and culture medium used. The tendency to form aggregates represents an important step for symbiosis.

The interactions are sufficient to form a dense texture similar to nativ thallus, known as soredial – like stage only after 90 or even 120 days for *Usnea barbata* species (Voicu D., Brezeanu A., 2008, Yamamoto *et al.*, 1985). A gradient density is useful for symbiosis establishment of powerful binding sites between symbionts.

Natural extracts supplemented media shortened the period of recognition and increased the number of interaction sites (fig. 3).

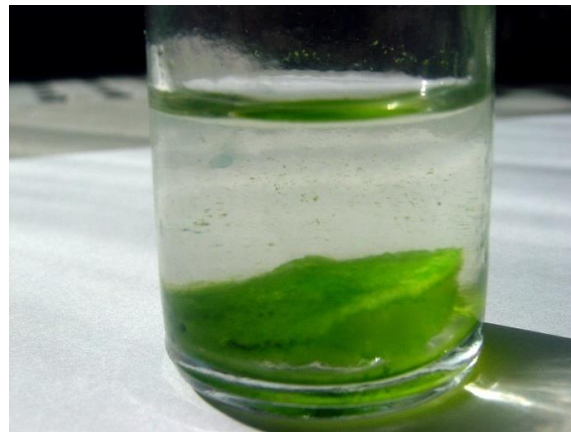


Fig. 3. *Usnea barbata* on kinetin (0.1 mg/l) enriched medium nine month from incubation – lichen radiate structure by photobiont dominated (Voicu D., Brezeanu A., 2008).

Mycobionts cultures took a long period of time (about one year), on MY medium (pH 5) at 15° C in the dark.

To avoid a degenerative phenomenon, subcultures were made in a range of 2 months.

Starting from the idea that cryopreservation is a time saving method to maintain the original variability of the germplasm for a long and very long time (from days to years and even hundreds of years) in independent conditions, we succeed to preserve for now, *Pseudevernia furfuracea* L., (Banciu and Cristian, 2015) in the sense that after thawing the explants maintained the viability.

Polyphenols concentration and antioxidant activity are preliminary results, revealed in Table 1 and comparative graphical representation (Fig. 10).

The main ultrastructural aspects of lichen symbionts were pointed (Voicu D., Toma N., Toma F.A., 2007; Brezeanu A., Voicu D., 2008). The interactions between them are materialised in new structures as a results of the culture conditions, like:

- an additional hydrophobic layer covering the mycobiont and photobiont cells.

– a neostructure, named – the interface (Fig. 5), between the mycobiont and phycobiont – cooperating the both partners to the biogenesis. By interface differentiating, the two partners cell walls lose their individuality, forming a tripartite structure. Beside this, mycobionts of the investigated lichens have a three-layered structure (Fig. 8).

Regarding the peculiarities of organelles, the chloroplast from “*in vitro*” aposymbiotically cultured photobionts present a primitive, lamellar, parallel-arranged system of thylakoids, with pyrenoglobules (Figs. 6, 7).

On the other hand, mycobionts from native thalli contain concentric bodies – a type of particular inclusion with important diagnostical feature; they are not found “*in vitro*” conditions. Mycobiont plasmallema is strongly folded. Paramural bodies are also specific for mycobionts from native thallus.

Some modifications “*in vitro*” conditions like the decrease in number and amplitude of invaginations and concentric bodies (Fig. 9) is another important aspect and also the enhanced number of mitochondria, suggesting a possible nutritional stress for *Cetraria islandica* mycobionts “*in vitro*” conditions.

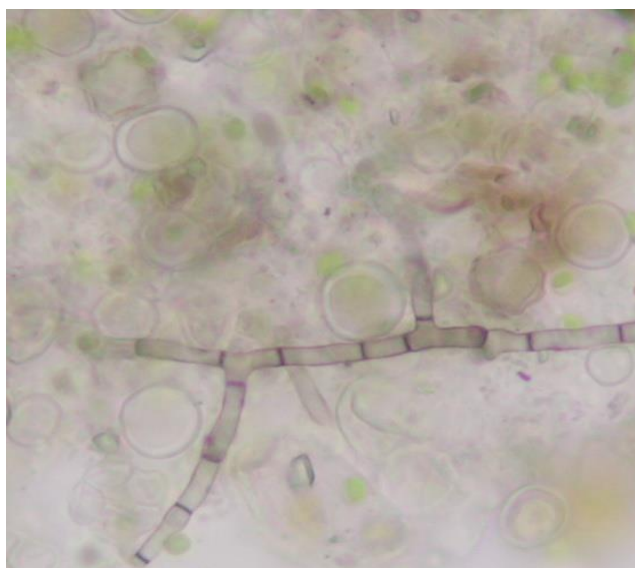


Fig. 4. Mycobiont – phycobiont interactions (oc. 10, ob. 40)
(Voicu D., Brezeanu A., 2008).

Table 1

Polyphenols concentration and antioxidant activity in methanol extracts
(Cristian D., Mitoi M., Brezeanu A., 2013)

Sample	Polyphenols ($\mu\text{g GAE/mg fresh weight}$)	Antioxidant activity ($\mu\text{g Trolox equivalents/mg fresh weight}$)
<i>In vitro</i> culture	21.022	90.529
Native thalli	99.296	194.758

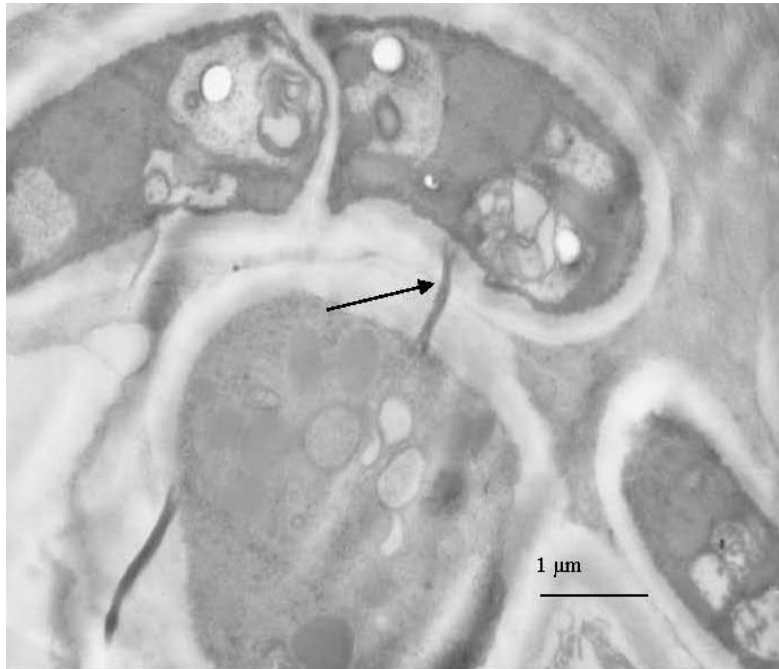


Fig. 5. Gelatinous parietal layer (see arrow) that illustrates the intimate connection between the mycobiont and phycobiont of *Usnea barbata*; phycobiont cell surrounded by mycobiont. The scale bar is 1 μm . (Brezeanu A., Voicu D., 2008).

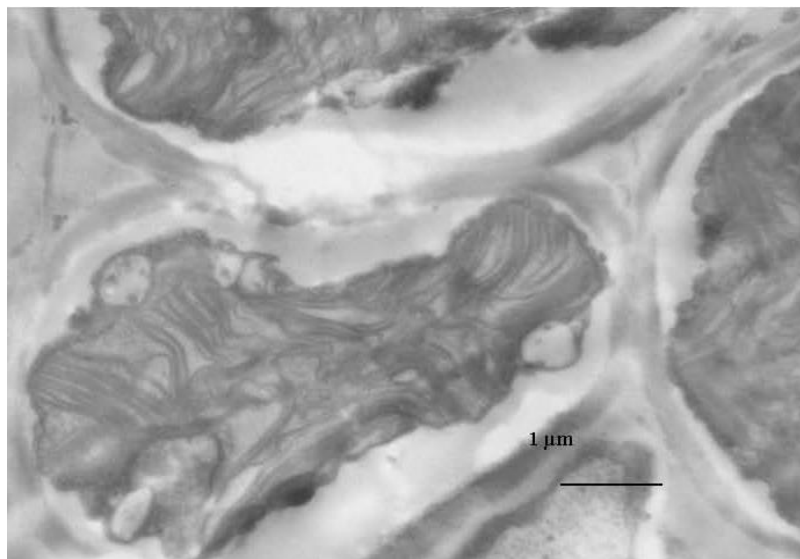


Fig. 6. Cross section through *U. barbata* phycobiont. Parallel arrangement of thylakoids is observed. The scale bar is 1 μm . (Brezeanu A., Voicu D., 2008).

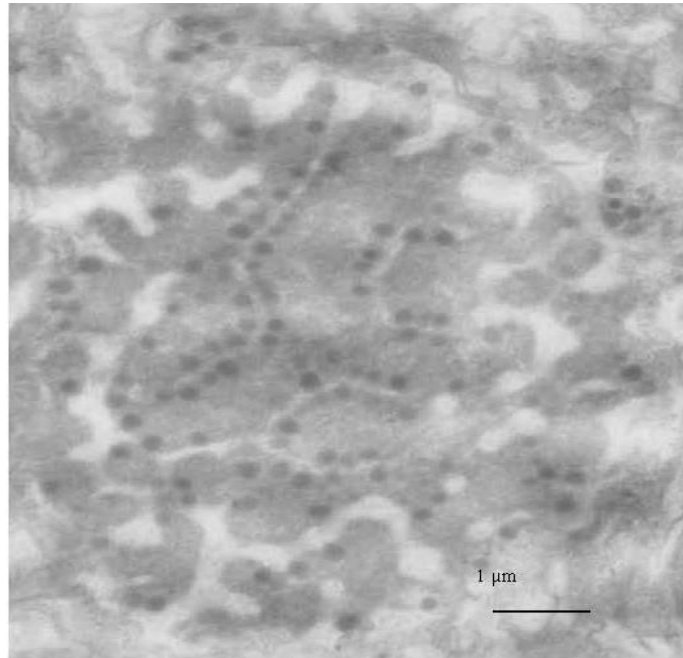


Fig. 7. Pyrenoglobules arranged along the chloroplast thylakoids (see arrangement). The scale bar is 1 μm . (Brezeanu A., Voicu D., 2008).

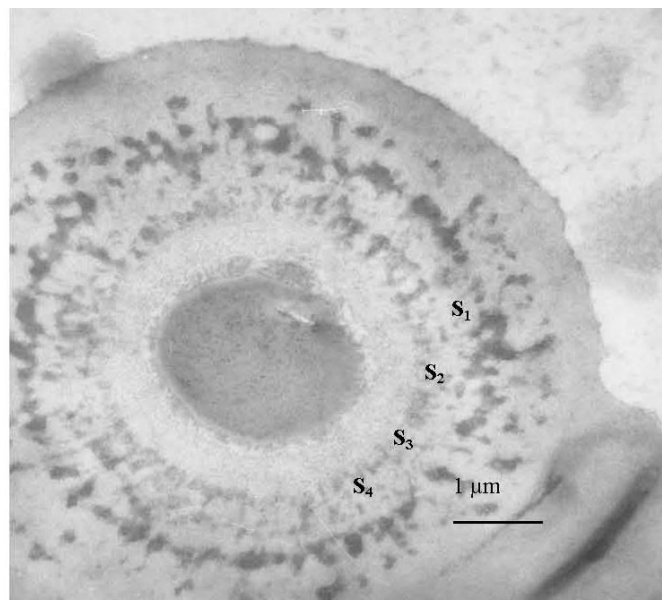


Fig. 8. Electronmicroscopical image of mycobiont cells in *U. barbata in vitro*. Three layered structure of cell wall. The scale bar is 1 μm . (Brezeanu A., Voicu D., 2008).

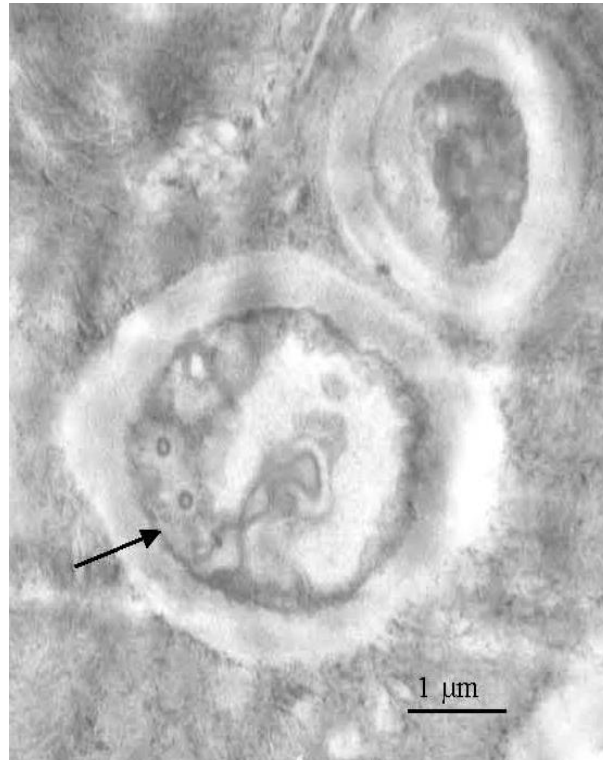


Fig. 9. Cross section through mycobiont cell. *In vitro*, concentric or ellipsoidal bodies are observed (see arrow). The scale bar is 1 μm . (Brezeanu A., Voicu D., 2008).

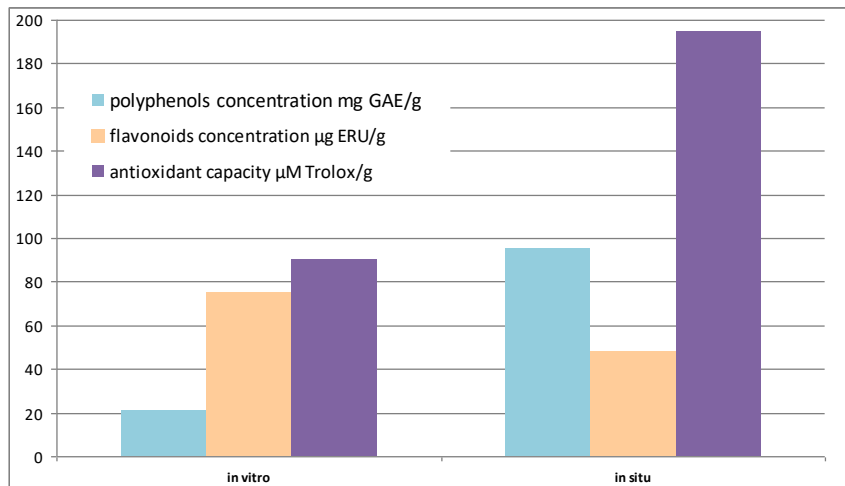


Fig. 10. Polyphenols concentration and antioxidant activity in methanol extracts *in vitro* and *in situ*.

CONCLUSIONS

Our studies regarding the possibility of using the “in vitro” systems for lichen species revealed merely the perspectives offered by them allowing the understanding of the cellular and molecular mechanisms involved in the establishment of this unique type of symbiosis, of cell recognition process and especially biotechnological, biosynthesis of bioactive compounds.

One noteworthy aspect was the highlighting of differentiated reactivity besides the “in vitro” conditions of lichen species analyzed, namely *Usnea barbata* (L.) Weber ex Wigg., *Cetraria islandica* (L.) Ach. and *Xanthoria parietina*, followed by *Pseudoevernia furfuracea*. We showed that the processes involved in “in vitro” cultures initiation and artificial resynthesis of symbiotrophic formations of lichen species are very complex, laborious, species specific.

So, every species requires special conditions both in term of type and size of the inoculum used for initiation, culture medium (composition, physical condition, hormones addition and incubation).

This is explained by the fact that each species presents morpho-anatomical and physiological different features and distinct ecological requirements.

Grinding paste methodology was useful for algae isolation and resynthesis experiments.

For mycobiont isolation, good results were obtained using apothecia like explants for *Xanthoria parietina* (L.) Th.Fr., *Usnea barbata* (L.) Weber ex Wigg, *Cetraria islandica* (L.) Ach.

Our findings showed that the methodology developed by us can be a promising alternative for lichen biomass obtaining of biotechnological interest and could be successfully adapted to other important species presenting a socio-economic interest.

Although they are not synthesized in great amount and the *in vitro* culture does not affect their biosynthesis, phenolic compounds are important for experimental systems use for that purpose.

Cryoconservation, frequently used for numerous higher plants species but not so frequently for lichens, proved to be a propitious method to maintain the cultures for long periods of time.

Also, our experiments allowed us to appreciate that the experimental system used by us specific for each species does not induce severe alterations on the inner structure and biosynthetic pathway of the cells and could represent a feasible alternative methodology for lichen biomass obtaining of biotechnological interest as well as for bioconservation of endemic or critically endangered species.

In perspective, for the future researches, we are interested in studying the elicitation involving the receptors from the mycobiont cell walls and the ligands from the photobiontic cell walls.

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POLLINATION ECOLOGY OF *RHYNCHOSIA SUAVEOLENS* (L.F.) DC. (FABACEAE), A PERENNIAL ERECT SHRUB IN THE SOUTHERN EASTERN GHATS, ANDHRA PRADESH, INDIA

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Abstract. *Rhynchosia suaveolens* is a perennial erect shrub. It flowers during October–January with peak flowering during November. The flowers are hermaphroditic, nectariferous, self-compatible and have an explosive pollination mechanism adapted for pollination by bees. They do not fruit through autonomous selfing but fruit through manipulated selfing, geitonogamy and xenogamy mediated by pollen vectoring bees. Lycaenid butterflies also visit, especially during peak flowering season, but they principally act as nectar robbers. The flowers not visited by bees fall off while those visited and pollinated by them set fruit. Seed dispersal occurs by explosive pod dehiscence. Perennial root stock produces new foliage during rainy season. Seeds also germinate at the same time but their continued growth is subject to the availability of soil moisture content. Therefore, *R. suaveolens* expands its population size in areas where soil is sufficiently moist and also in areas where trees or woody shrubs provide shade. Since the plant is not widely distributed, its occurrence is almost unknown to locals and hence is not used for any purpose.

Keywords: *Rhynchosia suaveolens*, hermaphroditism, explosive pollination mechanism, melittophilily, explosive pod dehiscence.

INTRODUCTION

Legume seeds are important sources of nutrients that meet our high quality dietary protein requirements (Perumal *et al.*, 2001; Escudero *et al.*, 2006). The under-utilized legumes, which have tremendous potential for commercial exploitation but remain ignored, offer a good scope in this context (Bhag Mal, 1992). Accounts of important under-exploited pulses which await exploration for food, fodder, energy and industrial purposes have been given (Siddhuraju *et al.*, 2000; Kalidass & Mohan, 2012). *Rhynchosia* is such a legume which is under-utilized despite its potential to provide high protein requirements. In this context, some work has been carried out on the biochemical and nutritional composition of *R. venulosa*, *R. hirta*, *R. cana*,

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R. filipes, *R. rufescens* and *R. suaveolens* by different workers (Lovelace, 1977; Murthy & Kandimalla, 2007; Kalidass & Mohan, 2012). Their work showed that these legumes are potential sources of protein, minerals and energy supplements for livestock and humans. The high protein value of *R. suaveolens* is in tune with its consumption by the tribes of Kani and Kannikars of southern Western Ghats of Tamil Nadu, India and hence it has potential for human nutrition (Kalidass & Mohan, 2012). This species has been reported to be distributed only in India and Sri Lanka (Manjunatha *et al.*, 2004). The knowledge of its reproductive ecology is essential to understand its sexual mode of reproduction and develop it as an edible legume crop. But, this information is totally lacking and hence the present study is contemplated to provide details of floral biology, pollination mechanism, pollinators and seed dispersal.

MATERIALS AND METHODS

STUDY SITE

The study region is an integral part of Southern Eastern Ghats of Andhra Pradesh in Peninsular India. The area is located at 13°40'N latitude and 79°19'E longitude. The exact study area is the forest cover of Tirumala Hills, a constituent of Seshachalam Hill Range in Chittoor District, Andhra Pradesh. The entire region represents the deciduous forest ecosystem. The site is characterized by a combination of rocky, undulating and steep terrain with some litter content formed from grass and other herbaceous plants. In this area, *Rhynchosia suaveolens* grows as scattered individuals or here and there as small populations in moist soils or shaded areas.

FLOWERING AND FLORAL BIOLOGY

Flowering season was defined based on regular field trips made. Twenty inflorescences were tagged and followed to record the length of flowering and the number of flowers produced. Anthesis was initially recorded by observing twenty five marked mature buds in the field. Later, the observations were repeated five times on different days in order to provide an accurate anthesis schedule. Similarly, the mature buds were followed for recording the time of anther dehiscence. The presentation pattern of pollen was also investigated by recording how anthers dehisced and confirmed by observing the anthers under a 10× hand lens. The details of flower morphology such as flower sex, shape, size, colour, odour, sepals, petals, stamens and ovary were described based on twenty five flowers randomly collected from five plants. Observations regarding the position and spatial relationships of stamens and stigma in mature bud, at anthesis and during the flower-life with reference to self and/or cross-pollination, were made very carefully.

POLLEN OUTPUT

Thirty mature but un-dehisced anthers from five different plants were collected and placed in a Petri dish. Later, each time a single anther was taken out and placed on a clean microscope slide (75 × 25 mm) and dabbed with a needle in a drop of lactophenol-aniline-blue. The anther tissue was then observed under the microscope for pollen, if any, and if pollen grains were not there, the tissue was removed from the slide. The pollen mass was drawn into a band, and the total number of pollen grains was counted under a compound microscope (40x objective, 10x eye piece). This procedure was followed for counting the number of pollen grains in each anther collected. Based on these counts, the mean number of pollen produced per anther was determined. The mean pollen output per anther was multiplied by the number of anthers in the flower for obtaining the mean number of pollen grains per flower. The characteristics of pollen grains were also recorded.

POLLEN-OVULE RATIO

The pollen-ovule ratio was determined by dividing the average number of pollen grains per flower by the number of ovules per flower. The value thus obtained was taken as pollen-ovule ratio (Cruden, 1977).

NECTAR CHARACTERS

The presence of nectar was determined by observing the mature buds and open flowers. The average volume of nectar per flower was determined and expressed in μL ; for this ten flowers were used. The flowers used for this purpose were bagged at mature bud stage, opened after cessation of nectar secretion and squeezed nectar into micropipette for measuring the volume of nectar. Nectar sugar concentration was determined using a Hand Sugar Refractometer (Erma, Japan). Ten samples were used for examining the range of sugar concentration in the nectar. For the analysis of sugar types, paper chromatography method described by Harborne (1973) was followed. Nectar was placed on Whatman No. 1 of filter paper along with standard samples of glucose, fructose and sucrose. The paper was run ascendingly for 24 hours with a solvent system of n-butanol-acetone-water (4:5:1), sprayed with aniline oxalate spray reagent and dried at 120°C in an electric oven for 20 minutes for the development of spots from the nectar and the standard sugars. Then, the sugar types present and also the most dominant sugar type were recorded based on the area and colour intensity of the spot. The sugar content/ flower is expressed as the product of nectar volume and sugar concentration per unit volume, $\text{mg}/\mu\text{L}$. This is done by first noting the conversion value for the recorded sugar concentration on the refractometer scale and then by multiplying it with the volume of nectar/flower. Table 5.6 given in Dafni *et al.* (2005) was followed for recording the conversion value to mg of sugars present in one μL of nectar.

Table 1
Anthesis as a function of time in *Rhynchosia suaveolens*

Time (h)	No. of flowers anthesed	Percentage of Anthesis
0800	0	0
0900	0	0
1000	13	17
1100	27	36
1200	26	35
1300	6	8
1400	3	4
1500	0	0

No. of mature buds tagged: 75

STIGMA RECEPTIVITY

In visual method, the stigma physical state (wet or dry) was considered to record the commencement of receptivity. H₂O₂ test as given in Dafni *et al.* (2005) was followed for the confirmation of stigma receptivity period.

BREEDING SYSTEMS

Mature flower buds of some inflorescences on different individuals were tagged and enclosed in paper bags. They were tested in the following way and the number of flower buds used for each mode of pollination was given in Table 2.

1. The flowers were fine-mesh bagged without hand pollination for autonomous autogamy.
2. The stigmas of flowers were pollinated with the pollen of the same flower manually by using a brush; they were bagged and followed to observe fruit set in manipulated autogamy.

Table 2
Results of breeding systems in *Rhynchosia suaveolens*

No. of flowers pollinated	No. of fruits formed	Fruit set (%)
35	0	0
50	9	18
50	34	68
50	41	82
853	523	61

3. The emasculated flowers were hand-pollinated with the pollen of a different flower on the same plant; they were bagged and followed for fruit set in geitonogamy.

4. The emasculated flowers were pollinated with the pollen of a different individual plant; they were bagged and followed for fruit set in xenogamy.

All these categories of flower pollinations were followed for fruit set. If fruit set is there, the percentage of fruit set was calculated for each mode.

FLOWER-VISITORS

The flower foragers included only bees and butterflies. The hourly foraging visits of insect species were recorded on 3 or 4 occasions depending on the possibility and the data was tabulated to use the same for further analysis. Fully blooming plants were selected to record the foraging visits of insects. The data obtained was used to calculate the percentage of foraging visits made by each species per day and also to calculate the percentage of foraging visits of each species per day in order to understand the relative importance of each species and category of insects. Their foraging behaviour was observed on a number of occasions for the mode of approach, landing, probing behaviour, the type of forage collected, contact with essential organs to result in pollination, inter-plant foraging activity in terms of cross-pollination.

DETERMINATION OF POLLEN CARRYOVER EFFICIENCY OF INSECT FORAGERS

Ten specimens of each insect species were captured from flowers and brought to the laboratory. Each specimen was washed first in ethyl alcohol and the contents stained with aniline-blue on a glass slide and observed under microscope to count the number of pollen grains present. From this, the average number of pollen grains carried by each species was calculated to know the pollen carryover efficiency of different species.

NATURAL FRUIT SET, SEED DISPERSAL AND SEEDLING ECOLOGY

A sample of flowers on twenty five plants were tagged on different plants prior to anthesis and followed for fruit set rate in open-pollinations. Fruit maturation period, fruit dehiscence and seed dispersal aspects were observed to the extent possible. Field observations were also made on fruit and seed dispersal modes, seed germination and seedling establishment to the extent possible.

PHOTOGRAPHY

Plant habitat, flowering inflorescences, and flower and fruit details were photographed with Nikon D40X Digital SLR (10.1 pixel) and TZ240 Stereo Zoom Microscope with SP-350 Olympus Digital Camera (8.1 pixel). Olympus Binoculars (PX35 DPSR Model) was also used to make field observations. Magnus Compound Microscope – 5×, 10×, 40× and 100× magnification was used for studying the pollen characteristics.

RESULTS

PHENOLOGY

It is a sweet-scented perennial erect hairy shrub with slender stem that grows moist, shaded areas (Figure 4a); it is also found to grow nearby streams where the soil is sufficiently moist. The plant re-grows from below ground perennial root stock and from the seed during June–September during which growth and leaf flushing occurs. The leaves are trifoliolate with reticulate venation. The leaflets are petiolate, ovate-acuminate, and puberulous, especially beneath. The flowering occurs during October–January with peak flowering in November (Figure 4b,c). The plants wither and disappear in April. The flower stalks arise in leaf-axils and each stalk is 2-flowered.

FLOWER MORPHOLOGY

The flowers are pedicellate, small (9.1 ± 0.5 mm long and 8.3 ± 0.4 mm wide), green with strong scarlet tinge, odorless, papilionaceous, zygomorphic and bisexual. The calyx is green with yellow tinge and consists of 5 free, lanceolate, acuminate, pubescent, 5.4 ± 0.6 mm long sepals. The corolla is red outside, yellow inside, pubescent, specialized and consists of upper standard petal, two wing petals and two keel petals. The standard petal is large (7.4 ± 0.6 mm long and 6.7 ± 0.4 mm wide), red lines inside at the bottom which serves as nectar guide; the petal base is clawed and consists of two inflexed fingernail auricles. The standard petal envelops the rest of the petals in bud but reflexes when the flower blooms. The two adjacent petals (7.3 ± 0.4 mm long and 2.8 ± 0.4 mm wide), called wing petals, surround the two bottom petals, called keel petals (7.2 ± 0.3 mm long and 2.8 ± 0.2 mm wide). The keel petals form a proximal cylindrical part and a distal part consisting of a pressed angular pouch, with an acute porate tip in which the stamens and stigma are housed. The keel and the wing petals are attached by means of two notched folds. The wing petals serve as a lighting platform for insects visiting the flowers. The stamens are ten, 5.7 ± 0.4 mm long, diadelphous; nine filaments are fused by the basal part into a sheath open along the upper side while the tenth filament is free and lies on the others (Figure 4i). The distal parts of the filaments are free and contain 1.2 ± 0.3 mm long uniform ditheous anthers. The ovary is sessile, green, villous, 2.5 ± 0.5 mm long (Figure 4l) and lies in the sheath of the filaments along the cylindrical part of the keel. It is monocarpellary and monolocular with two ovules arranged on marginal placentation (Figure 4n). It has a long filiform style with a capitate wet shiny stigma, both together account for a length of 6.5 ± 0.6 mm (Figure 4m). The stigma is situated slightly beyond the length of the anthers. The distal portion of free filaments and style and stigma are incurved and clamped into the keel petals.

FLORAL BIOLOGY

Mature buds open during 1000–1400 h with peak anthesis during 1100–1200 h (Table 1). Unfolding of the standard petal and wing petals indicates flowering opening. The keel petals do not unfold and remain in their original position as in mature bud stage (Figure 4d–h). All the ten anthers in a flower dehisce at the same time by longitudinal slits in mature bud stage (Figure 4j). The number of pollen grains per anther is 773.4 ± 42.34 and per flower is 7,734. The pollen-ovule ratio is 3,867:1. The pollen grains are monads, spheroidal, 24.86 ± 5.24 μm in size, powdery and tricolporate, angulaperturate with reticulate exine (Figure 4k). A nectariferous disc is present at the base of the ovary. The initiation of nectar secretion occurs during mature bud stage and its cessation occurs an hour after anthesis. Individual flowers produce 1.4 ± 0.21 μL of nectar with 0.45 mg of sugar. The nectar sugar concentration is 29% (Range 27–31%) consisting of sucrose, glucose and fructose with the first as dominant. Nectar is deeply concealed and it is open through two windows between the joined and the free filaments at the flower base. These windows allow access to the nectar. The stigma attains receptivity during anthesis and remains receptive for about three hours. After three hours of anthesis, the standard, wing and keel petals gradually move close to each other enclosing the reproductive organs. The closed flowers remain so even during most part of the fruit development. The calyx initially encloses the ovary and subsequently turns light brown and discloses the ovary since the latter gradually bulges and develops into a seeded pod.

POLLINATION MECHANISM

The reproductive column is held under pressure within the keel part in open flowers and it is exposed when the pollinator presses against the wing and the keel petals. When insects land on the wing petals, the latter causes the keel petals to release the reproductive column explosively. Consequently, the reproductive column snaps forward against the standard petal causing most of the pollen to be instantly released and the pollen thus released comes into contact with the ventral side of the insect body. Since the incurved stigma is situated above the height of the anthers, it strikes the insect body first due to which cross-pollination occurs if the insect visited the other flowers previously and carried pollen on its ventral side and also then the pollen ejected from the anthers powders the ventral side of the insect instantly. If it is the first visit for the insect to the flower, then it effects self-pollination upon explosive release of reproductive column from the keel boat. With the departure of the insect from the flower, the reproductive column does not return back to its former position but the keel moves forward partly covering the stamens and stigma. The downward movement of keel petals occurs in each subsequent foraging visits by appropriate insects. Tripping of keel boat can also occur due to heavy rain or high temperature that weaken turgidity of the restraining keel tissues.

But, the tripping due to these two factors is ruled out since the plant flowers during winter season when heavy rains do not normally occur and the temperature usually stands low. If the flower is untouched or tripping to keel did not occur, the reproductive column is never exposed and remains enclosed in the keel boat. Such flowers fall off subsequently upon withering without fruit set.

BREEDING SYSTEMS

In mature buds, anthers dehisce but autonomous autogamy does not occur. Fruit set is absent in un-manipulated autogamy, 18% in hand-pollinated autogamy, 68% in geitonogamy, 82% in xenogamy and 61% in open-pollination (Table 3).

Table 3

Natural fruit/seed set rate in *Rhynchosia suaveolens*

No. of flowers sampled	No. of flowers set fruit	Fruit set (%)	Seed set (%)	Rate of seed set/fruit					
				No. of 1-seeded fruits	1-seeded fruit set (%)	No. of 2-seeded fruits	2-seeded fruit set (%)	No. of seedless fruits	Seedless fruits (%)
853	523	61	58	42	8	476	91	5	1

BEE POLLINATORS AND POLLINATION

The flowers were exclusively foraged by bees for pollen and nectar and by butterflies for nectar. The bees showed foraging activity during 1040–1630 h with peak activity during 1230–1330 h (Figure 1). The bees included *Apis florea* (Figure 5a), *Xylocopa latipes* (Figure 5b), *Ceratina* sp. (Figure 5c–f) and *Nomia* sp. (Figure 5g) (Apidae) (Table 4); they were regular and consistent foragers throughout the flowering season. The butterflies included only Lycaenids, *Castalius rosimon* (Figure 5h), *Freyeria trochylus*, *Lampides boeticus* and *Chilades pandava* (Figure 5i) (Table 4); they foraged during 1130–1530 h with peak activity during 1230–1330 h (Figure 2). Bee foraging visits constituted 84% and butterflies 16% of total foraging visits made in a given day during peak flowering phase (Figure 3). All bee species were regular and consistent foragers throughout the flowering season. In case of butterflies, they were occasional in their visits during initial and final phase of flowering while they were regular and consistent during peak phase of flowering. The bees were quite effective to trip the flowers while foraging for nectar and effect pollination. The bees landed on the wing petals and the keel, with their head near the standard. They then exerted a certain pressure with legs on the wing petals until these and the keel bent downwards, and then proceeded to collect nectar during which the bee's abdomen appeared pollen smothered (sternotribic pollen deposition). To collect pollen, the bees took "U" turn after nectar collection and proceeded to the stamens to collect pollen. The butterflies were in-effective to trip the flowers but foraged for nectar from both un-tripped as well as tripped flowers. In case of un-tripped

flowers, they slowly inserted their proboscis into the base of standard petal to collect nectar due to which there was no tripping of the keel petals at all. In this case, they did not effect pollination but acted as only nectar robbers. In case of tripped flowers, they exhibited the same probing behavior for nectar collection but the contact between their proboscis or other body parts and the stamens and stigma took place only when they probed the flower from the front. When they probed the flower from the sides of the flower, there was no contact between their proboscis and the stamens and stigma. From this, it was found that the butterflies acted as pollinators when they probed the flowers from the front and as nectar robbers when they probed the flowers from the sides. The bees and butterflies were found to visit the same flowers on the same and different plants several times during the day and such a foraging activity was considered to be promoting cross-pollination rate. The body washings of bees and butterflies showed that they carried pollen to different extents; the bees showed the pollen carrying capacity ranging from 321 to 1793 while the butterflies, especially on their proboscis carried pollen ranging from 26 to 89 (Table 5). These data indicated that both bees and butterflies are pollen carriers and effect pollination to different extents.

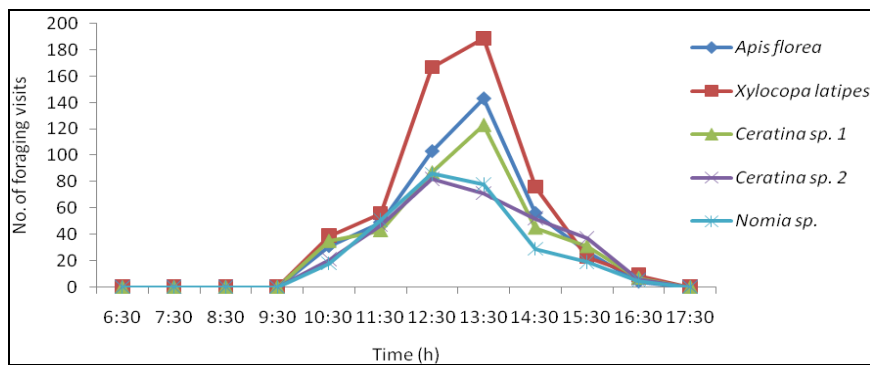


Fig. 1. Hourly foraging activity of bees on *Rhynchosia suaveolens*.

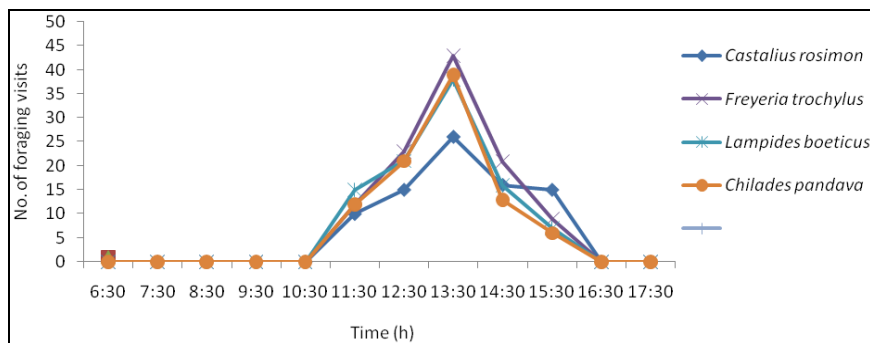


Fig. 2. Hourly foraging activity of butterflies on *Rhynchosia suaveolens*.

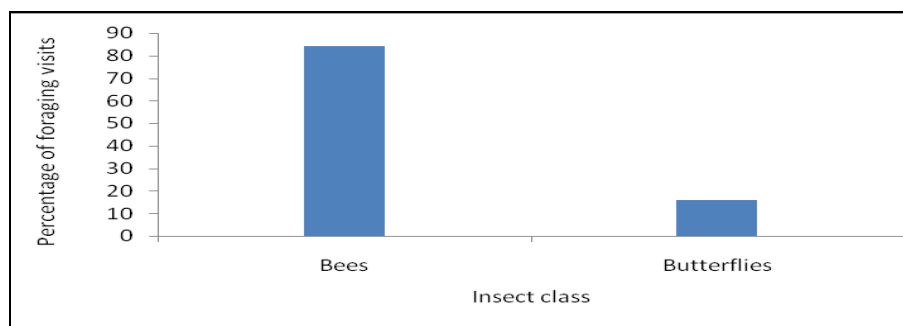


Fig. 3. Percentage of foraging visits of bees and butterflies on *Rhynchosia suaveolens*.

Table 4

List of insect foragers on *Rhynchosia suaveolens*

Order	Family	Sub-family	Genus	Species	Common Name	Foraging schedule	Forage collected
Hymenoptera	Apidae	Apinae	<i>Apis</i>	<i>florea</i>	Dwarf bee	1030–1630	Nectar Pollen
				<i>latipes</i> Drury	Large carpenter bee	1030–1630	Nectar
		Xylocopinae	<i>Ceratina</i>	sp. 1	Small carpenter bee	1030–1630	Nectar Pollen
				sp. 2	Small carpenter bee	1030–1630	Nectar Pollen
				sp.	Alkali bee	1030–1630	Nectar Pollen
		Lepidoptera	Lycaenidae	Polyommatae	<i>Castalius</i>	<i>rosimon</i> F.	The Common Pierrot
<i>trochylus</i> Freyer	Grass Jewel					1130–1530	Nectar
<i>Lampides</i>	<i>boeticus</i> L.				The Pea Blue	1130–1530	Nectar
	<i>pandava</i> Horsfield				The Plains Cupid	1130–1530	Nectar

Table 5

Pollen recorded in the body washings of bee foragers on *Rhynchosia suaveolens*

Insect species	Sample size (N)	Number of pollen grains		
		Range	Mean	S.D
<i>Apis florea</i>	10	423–878	631.1	153.06
<i>Xylocopa latipes</i>	10	762–1793	1325.9	381.41
<i>Ceratina</i> sp. 1	10	378–654	501.9	104.10
<i>Ceratina</i> sp. 2	10	398–691	544.7	110.41
<i>Nomia</i> sp.	10	321–586	452.1	91.87
<i>Castalius rosimon</i>	10	62–89	75.8	8.59
<i>Freyeria trochylus</i>	10	43–78	62.1	12.14
<i>Lampides boeticus</i>	10	27–61	48.1	12.40
<i>Chilades pandava</i>	10	26–56	33.7	8.58

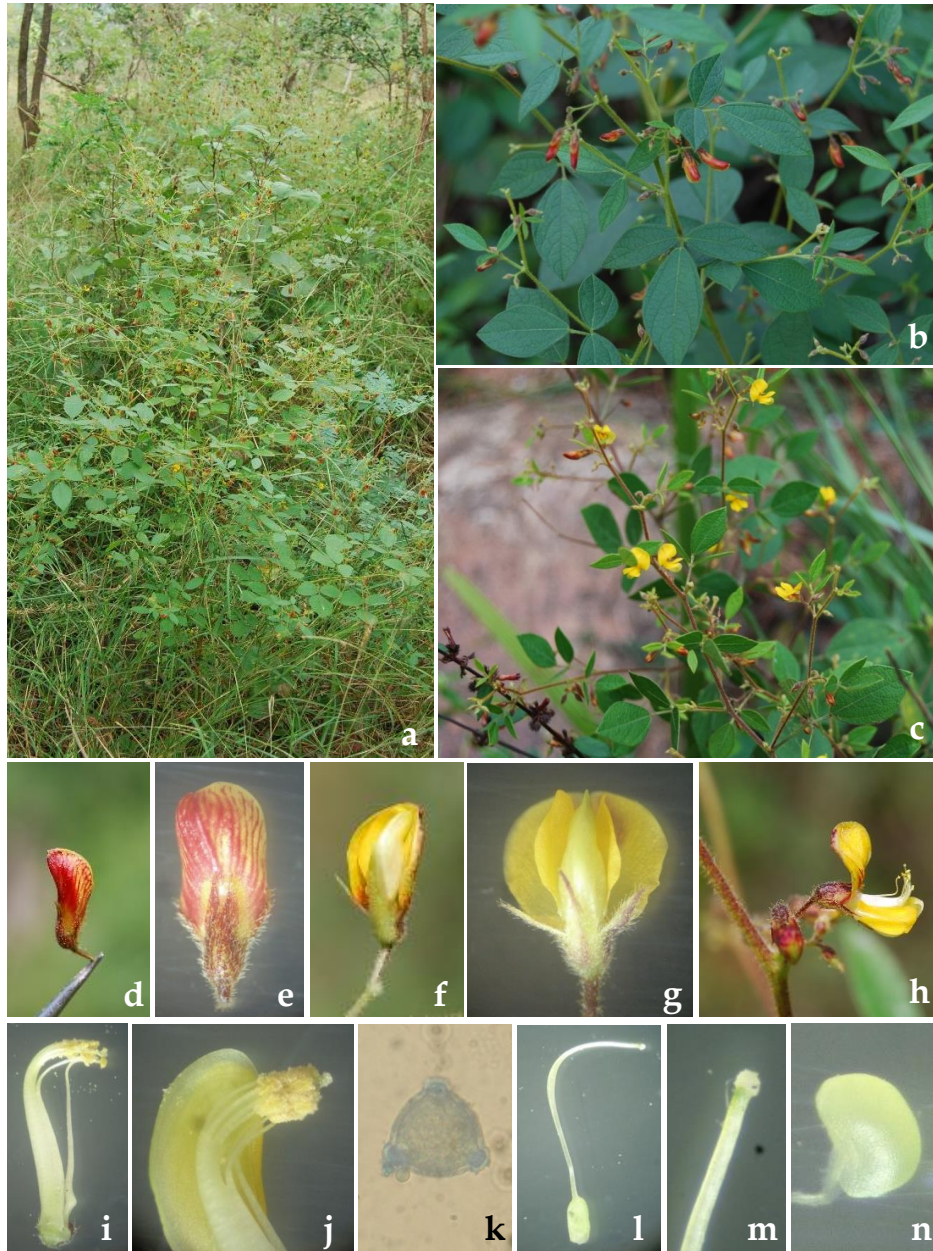


Fig. 4. *Rhynchosia suaveolens*: a. Habitat, b. Budding phase, c. Flowering, d–g. Different stages of anthesis, h. Keel and wing petals with stamens and stigma, i. Diadelphous stamens, j. Dehisced anthers, k. Pollen grain, l. Gynoecium, m. Capitate stigma, n. ovule.

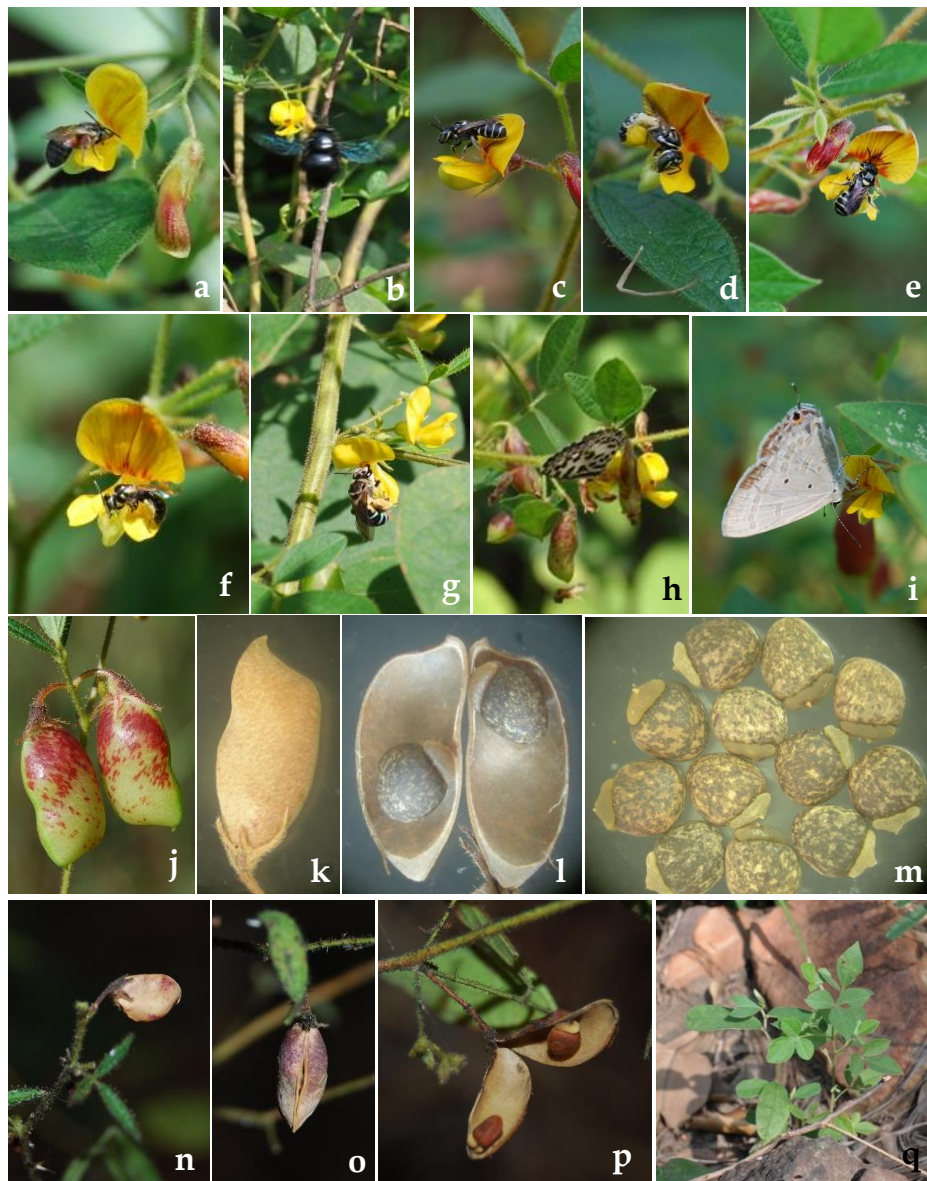


Fig. 5. *Rhynchosia suaveolens*: a. *Apis florea* collecting nectar, b. *Xylocopa latipes* collecting nectar, c. & d. *Ceratina* sp. 1 collecting pollen, e. *Ceratina* sp. 1 collecting nectar, f. *Ceratina* sp. 2 collecting pollen, g. *Nomia* sp. collecting nectar, h–i. *Lycaenids* – h. *Castalius rosimon* collecting nectar, i. *Chilades pandava* collecting nectar, j. Maturing pods, k. Mature, dry pod, l. 2-seeded pod, m. Seeds, n–p. Explosive pod dehiscence, q. Seedling.

FRUITING BEHAVIOR

The fruit growth and development begins immediately after pollination and fertilization. The fruits mature within three weeks. The sepals enclose the growing fruit initially and the fruit emerges out of the sepals gradually with its gradual growth and development. Fruit is green initially with scarlet red tinge and brown when ripe and dry (Figure 5 j,k). It is a non-fleshy, hairy, oblong, 11.2 ± 1.5 mm long, 5.1 ± 0.4 mm wide, compressed, rounded and apiculated pod. The pods produced mostly two seeds (Figure 5l) while a few produce one seed. Fruit set without seeds also exists but it is negligible. The fruit set rate in open-pollination mode is 61% out of which 2-seeded pod set rate is 91%, 1-seeded pod set rate is 8% and seedless pod set rate is 1%. Total seed set rate is 58% (Table 3).

SEED ECOLOGY

Mature and dry fruits display explosive dehiscence to disperse seeds (Figure 5n-p). The pod with bi-valvate configuration dehisce to eject the seed. The seed is greyish brown, striate, reniform, finely pubescent, 4.6 ± 0.4 mm long, 3.6 ± 0.3 mm wide, shiny and strophiolate (Figure 5m). Seeds germinate during rainy season which starts from June onwards. Seedlings grow continually but their growth rate is subject to the availability of moisture status of the soil (Figure 5q). In areas where soil is sufficiently moist, seeds continue growth and produce mature plants within two months and subsequently commence flowering and fruiting.

DISCUSSION

Distribution range of *Rhynchosia suaveolens* is largely unknown as there is almost no information regarding its distribution and habitats of occurrence. Manjunatha *et al.* (2004) mentioned that it is distributed in India and Sri Lanka only. The present study shows that *R. suaveolens* occurs in moist soils or shaded areas in dry deciduous forest areas of Eastern Ghats of Andhra Pradesh. It usually occurs as scattered individuals or as small populations here and there. As a perennial erect shrub, it grows from seed as well as from perennial root stock during rainy season. The flowering season is well defined and is confined to winter season. Individual plants produce a small number of flowers during their lifetime due to production of a few inflorescences and each of which produces two flowers. Individual flowers are prominent only by their red to yellow coloured corolla but not by their position since they occur within the foliage.

In *R. suaveolens*, hermaphroditic sexual system is functional due to production of fertile pollen grains and functional ovary. The flowers display the near synchronous hermaphroditism or homogamy due to the occurrence of anther dehiscence in

mature bud stage and receptivity of stigma during anthesis. The entire reproductive column stays inside the keel petals even after anthesis; in this situation, there is a likelihood of the occurrence autonomous autogamy. But, hand-pollination tests indicated that autonomous autogamy does not occur despite self-compatibility but it is functional when this mode of pollination is manipulated by brushing the stigma with its own pollen. Such a situation suggests that the flowers are essentially dependent on flower foragers for fruit set through self – as well as cross-pollination. It appears that the stigma although receptive blocks the germination of the self-pollen while it is in keel petals and hence, it essentially requires the rupture of its surface by a pollinator to allow the self – or cross pollen to germinate. Such a stigmatic regulatory function appears to have evolved to discourage selfing and promote out-crossing. Shivanna & Owens (1989) stated that the rupture of the stigmatic surface by pollinator permits the pollen to germinate in the flowers of Phaseoleae members with thick stigmatic cuticle. On the contrary, Castro & Agullo (1998) reported that in *Vigna*, a member of the tribe Phaseoleae, autonomous self-pollination may occur by spontaneous rupture of the stigmatic membrane. Similar stigmatic surface that prevents self-fertilization has also been reported in *Vicia faba* (tribe Viciaeae) (Lord & Heslop-Harrison, 1984) and in *Medicago scutellata* (tribe Trifolieae) (Krietner & Sorensen, 1985); however, in these species auto-fertile lines have been reported to have thin stigmatic cuticles allowing spontaneous disruption and self-fertilization. In *R. suaveolens*, the stigmatic surface appears to have thick cuticle and does not have the mechanism of causing spontaneous rupture to facilitate autonomous self-pollination. In effect, the tripping of keel petals appears to be essential to cause rupture on the stigmatic surface by the tripping agent due to which there is more likelihood of the occurrence of either geitonogamy or xenogamy. The fruit set rates recorded in hand-pollinated geitonogamy and xenogamy also substantiate that the plant is facultative xenogamous, a breeding system that is flexible and keeps the options open for both selfing and out-crossing mediated by pollen vectors.

Schrire (1989) stated that the ecological and evolutionary success of Leguminosae has been related to biotic pollination mechanisms. The three sub-families within this family have achieved a characteristic floral architecture, in which plants within the sub-family Papilionoideae have developed the most complex floral mechanisms. Plants within the Papilionoideae have zygomorphic flowers that are mainly bee-pollinated (Westerkamp, 1997); although bird pollination and bat pollination have also been recorded within the subfamily (Ortega-Olivencia *et al.*, 2005). In bee-pollinated flowers of Papilionoideae, it is assumed that each part of the corolla is specialized for a particular role in pollinator attraction and the success of pollination. The flag or standard petal attracts pollinators; the keel protects androecium and gynoecium and, together with the wings, provides a platform for the insects to land on. The wings also operate as levers that raise or lower the keel (Stirton, 1981). The flowers typical of pollination by the bee family Apidae are zygomorphic, bright yellow or blue with nectar guides, and frequently with hidden rewards such as those in the Lamiaceae, Scrophulariaceae, Fabaceae

and Orchidaceae (Faegri & van der Pijl, 1979). In the present study, the Fabaceae member, *R. suaveolens* has papilionaceous corolla with flag, wing and keel petals; the flag petal serves as a visual attractant, wing petals provide landing platform and keel petals protect the entire reproductive column. The flowers are typical of pollination by bees since they are zygomorphic, standard petal with nectar guide, hidden nectar at the corolla base and hidden pollen in keel petals.

Within the sub-family Papilionoideae, primary and secondary pollen presentations have been reported. In plants with primary pollen presentation, pollen is delivered directly from the anthers to the vector's body. In plants with secondary pollen presentation, pollen grains are delivered first on a floral part such as the keel petals in Papilionoideae and then on the body of the vector implying an accurate delivery of pollen on the vector's body (Howell *et al.*, 1993). These two pollen presentation patterns are associated with the four types of basic pollination mechanisms – valvular, pump, explosive and brush, all of them are associated with a particular floral architecture and kinetics. In the valvular type, pollen presentation is primary, whereas in the other three mechanisms, it is secondary (Yeo, 1993). In the explosive mechanism, commonly only one pollination event occurs and it has evolved independently in several tribes (Small, 1988), while in the other three mechanisms, repeated visitation is possible (Westerkamp, 1997). In the present study, *R. suaveolens* flowers have explosive pollination mechanism and deliver pollen directly from the anthers to the bee's body when keel petals are tripped by the foraging bee; this type pollen delivery is the representative of primary pollen presentation associated with explosive pollination mechanism. In the flowers, the staminal column is held under pressure within the keel, and when the tension is released by the forager, the same column snaps forward against the standard petal causing all the pollen to be instantly released. The reproductive column remains exposed and does not return back to its original state but the keel petals return back partially covering the stamens and stigma. The efficiency of explosive pollination mechanism depends on the ambient weather conditions, especially temperature and relative humidity. Since *R. suaveolens* flowers during winter season, it accordingly commences anthesis from late morning onwards by which time the ambient air will be relatively dry and hence is conducive for the efficient functioning of the explosive pollination mechanism. Further, the bees also commence their foraging activity as soon as the flowers are open and continue forage collection until the flowers close back. The concealment of the stamens within the keel petals until it is tripped is an advantage for the plant to secure pollen from unusual rains and ambient moisture conditions during the flowering season of this plant (Peter *et al.*, 2004).

Percival (1961) stated that plants with deep-tubed flowers tend to produce sucrose-rich nectar, whereas those with open or shallow-tubed flowers tend to be hexose-rich. Baker & Baker (1983) stated that flowers with long corolla tube possess more sucrose in their nectar while those with short tubes possess more hexoses in their nectar. In the present study, *R. suaveolens* with short corolla tube presents sucrose-rich nectar because the nectar is perfectly concealed and hence is

not exposed for the breakdown of sucrose into hexoses. Concealment of nectar in this species is adaptive to protect against microorganisms, particularly yeasts, whose metabolic activities dramatically change nectar chemistry and the plant gains a benefit from keeping the nectar as sterile as possible to maintain control over its chemical composition in order to maximize pollination rate by attracting appropriate pollinators (Herrera *et al.*, 2008). Honey bees prefer the flowers with sucrose as chief constituent of nectar (Kevan, 1995). The flowers pollinated by long-tongued bees produce sucrose-rich nectar (Baker & Baker, 1990). In line with this, *R. suaveolens* with melittophilous pollination syndrome also produces sucrose-rich nectar which is utilized exclusively by long-tongued bees. *Apis*, *Ceratina*, *Nomia* and *Megachile* bees recorded on this herb have been documented as long-tongued bees (Cruden *et al.*, 1983; Roubik, 1992; Roubik, 2006). Bee-flowers tend to produce small volume of nectar with higher sugar concentration than the nectar of flowers pollinated by other animals (Opler, 1983; Cruden *et al.*, 1983). Honey bees prefer sugar concentration of 20 to 40% in the nectar (Waller, 1972). On the contrary, Baker & Baker (1983) noted that honey bees prefer sugar concentration of 30 and 50% in the nectar. The honey bees have the ability to regurgitate liquid onto concentrated or even crystallized nectar, in this way, reduce its concentration so that it may be imbibed. The preferred sugar concentrations of nectar by other categories of bees have not been found in the literature. But, Pyke & Waser (1981) stated that the nectar sugar concentration of flowers pollinated by bees is generally higher than that of those pollinated by butterflies and hummingbirds; bee-pollinated flowers tend to produce nectar with sugar concentration more than 35% while butterfly or hummingbird pollinated flowers tend to produce nectar with sugar concentration ranged between 20 and 25%. In line with these reports, the present study shows that the flowers of *R. suaveolens* produce a small volume of nectar with 29% sugar concentration. Further, the energy yield from nectar appears to be in tune with the requirement of energy by bees. Therefore, *R. suaveolens* flowers with explosive pollination mechanism, primary pollen presentation, and hidden nectar and pollen have evolved to discourage other foragers from visiting the flowers and to ensure that the bees get the floral rewards. Accordingly, the flowers were pollinated principally by bees. Apart from bees, lycaenid butterflies also visit the flowers for nectar especially during peak flowering season but their probing behavior indicates that they do not cause tripping of flowers to result in pollination but collect nectar by slowly inserting their proboscis into the floral base. They collect nectar from tripped flowers with great ease with or without any contact between their proboscis and floral sex organs. Therefore, butterflies are principally nectar robbers and in effect, standing crop of nectar gets reduced. In this situation, the pollinating bees make multiple visits to the same flowers in quest of nectar and enhance pollination rate. Therefore, butterfly foraging activity is indirectly contributing to the enhanced natural fruit set rate. In line with this, natural fruit set rate recorded for *R. suaveolens* is the highest.

Cruden (1977) used the pollen-ovule (P/O) ratios as indicators of breeding systems of plants. He provided P/O ratios for different breeding systems – 168.5 +

22.1 for facultative autogamy, 798.6 + 87.7 for facultative xenogamy and 5859.2 + 936.5 for xenogamy. Several workers followed these P/O ratios to classify breeding systems of the plant species studied by them. Arroyo (1981) stated that the P/O varies according to the pollination mechanism within Papilionoideae. These authors suggested that the plants with explosive mechanism have a low P/O because a single pollinator visit is needed for efficient transference of pollen; this low P/O is a consequence of the highly specialized, irreversible pollination mechanism, which allows only one effective exchange of pollen with pollinators. Small (1988) stated that *Medicago* species of the tribe Trifolieae with explosive pollination mechanism displays the lowest pollen-ovule ratios. Lopez *et al.* (1999) recorded an explosive pollination mechanism with the highest pollen-ovule ratios in certain genera of the Fabaceae such as *Cytisus*, *Pterospartum*, *Teline*, *Ulex*, *Stauracanthus* and *Cytisophyllum*. Etcheverry *et al.* (2011) stated that the Fabaceae plants which they studied with an explosive pollination mechanism had intermediate pollen-ovule ratios. These authors mentioned that *Rhynchosia edulis* and *R. senna var. texana* have a valvular pollination mechanism with primary pollen presentation. Both the species are classified as obligate xenogamous based on P/O ratio but *R. edulis* has been found to be facultative xenogamous in hand-pollination tests. Craufurd & Prins (1979) reported that *R. sublobata* is self-compatible and facultative xenogamous in hand-pollination tests; it is pollinated by *Xylocopa* bees. In the present study, *R. suaveolens* shows the highest P/O ratio when compared to that of facultative xenogamy used by Cruden (1977). The highest P/O ratio in this plant species appears to be a consequence of pollen collection activity by bees. Therefore, it is inevitable for *R. suaveolens* to produce high P/O to compensate the pollen loss caused by pollen collectors and ensure the function of its vector-dependent facultative xenogamous breeding system.

Tran & Cavanagh (1984) reported that in Leguminosae, seeds of many taxa exhibit physical dormancy due to the presence of a water impermeable seed coat. With this dormancy, they remain viable for a long period of time. Ali *et al.* (2012) reported such physical dormancy in *Rhynchosia capitata* due to which this species is successful as a weed. Shaukat & Burhan (2000) described seed characteristics and the factors regulating germination of *R. minima* in Pakistan; it exhibits differential success in different habitats with different micro-climates. Rangaswamy and Nandakumar (1985) reported that the seed coat of *Rhynchosia minima* is composed of three gradative barriers to water uptake – a surface deposit of waxy material interfused with a lipoidal substance, β -sitosterol; a subjacent 3- μ m adcrustation of hemicellulose-cellulose complex; and a layer of palisade cells in which the secondary walls are impregnated with arabinan and the lumen contains tannin and phenolic compounds. The micropyle and hilum function as hygroscopic valves when seed coat breaks open. In *R. suaveolens*, seeds exhibit physical dormancy during dry season and respond to rainfall during wet season. Many seeds germinate in the vicinity of or away from the parental plants during rainy season but their continued growth is related to soil moisture and nutrient status. Since the plant is a perennial, its underground root stock also produces new growth during rainy season, flowers

and fruits during winter season. In *R. suaveolens*, the explosive pod dehiscence does not spread the seeds far away from parental sites but rain water causes the seeds to disperse to new places. Therefore, *R. suaveolens* is not a very widespread species and is primarily an inhabitant of moist soils or shaded areas, the preference of which makes it a typical mesophyte.

R. suaveolens is traditionally used as food by the tribes of Kani and kannikars of southern Western Ghats of Tamil Nadu, India (Kalidass & Mohan, 2012). A study on the biochemical and nutritional composition of this species showed it is an important source of protein, minerals and energy supplements for livestock and humans (Lovelace, 1977; Murthy & Kandimalla, 2007). Keeping this in view, further studies are suggested to evaluate the potentiality of this plant as a commercial legume crop to meet a part of nutritional requirements of livestock as well as humans.

Remanandan (1981) stated that *Rhynchosia*, being closely related to the genus *Cajanus*, some of its species can be used to provide substantial contributions towards crop improvement in pigeon pea. Furthermore, some species of *Rhynchosia* have been experimented in India to provide physiological resistance against insect pests such as pod-borer and pod-fly in pigeon pea. In this study, the seeds of *R. suaveolens* have not been infested by any pod-borer or pod-fly both in water-saturated and water-stress habitats suggesting that it has physiological resistance against insect pests. Therefore, intensive and extensive research is suggested to identify and select desirable genotypes of *R. suaveolens* that give physiological resistance against pod or seed pests in order to use them for crop improvement in pigeon pea or other legumes.

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EFFECT OF SODIUM AZIDE ON MORPHOLOGICAL CHARACTERS OF THREE TOMATO ACCESSIONS (*SOLANUM LYCOPERSICON* L.)

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Abstract. The effect of chemical mutagen (Sodium azide) was used to study the morphological characters of three accessions of tomato (*Solanum lycopersicon*) and to estimate the heritability of treated plants. Dry seeds of one accession (NGB01302) were collected from National Centre for Genetic Resources (NACGRAB) and two accessions (Tropimech and Tima) were collected from National Horticultural Research Institute (NIGHORT), Ibadan, Nigeria. The seeds were presoaked in water to test for viability. The seeds were soaked in sodium azide of concentrations 1mM, 2mM and 4mM in phosphate buffer of pH 4 for 4 hours and rinsed under a running tap to remove excess mutagen. The result of the study showed that the control of Tropimech had the highest germination (100%) while the lowest was recorded in treatment with 1mM. At four weeks of planting (4WAP), there was no significant difference between the heights of treated seedlings of Tropimech while the untreated seedlings were significantly different from the treated seeds. No significant difference was observed in both treated and untreated seedlings of Tima and NGB01302. Highly significant differences ($P < 0.05$) were observed in the number of leaves, number of branches, number of flowers, number of fruit and fruit weight in the three accessions. The heritability of the yield parameters (number of flower and fruit weight) were moderate and the genetic advance was high in the number of leaves, number of flowers and fruit weight. There was general reduction in all parameters under study with increasing concentration of sodium azide except in NGB01302 which showed an increase.

Keywords: Sodium azide, mutagen, *Solanum lycopersicon*, accession.

INTRODUCTION

Chemical mutagenesis is regarded as an effective and important tool in improving the yield and quality characters of crop plants. Sodium azide has also proved its worth as a chemical mutagen used to induce genetic variability (Srivastava *et al.*, 2011). Successful mutant isolation largely relies on the use of efficient mutagens. The role of mutation breeding in increasing the variability for quantitative traits in various crop plants has been proved beyond doubt by a

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number of scientists (Khan *et al.*, 1998; Das and Chakraborty, 1998; Rachovska and Dimova, 2000; Erdem and Oldacay, 2004; Khan and Goyal, 2009; Adamu and Aliyu, 2007; Siddiquis *et al.*, 2007; Nilan *et al.*, 1977).

Mutation induction offers the possibility of inducing desired attributes that either cannot be found in nature or have been lost during evaluation. Treatment of plant with mutagens alters genes or breaks chromosomes.

The main advantage of mutation breeding is the possibility of improving one or two characters without changing the genotype. Induced mutation has great potential and serves as a complimentary approach in the genetic improvement of crops (Novak *et al.*, 1992).

Tomato (*Solanum lycopersicon*) is an economically important plant, therefore the need to improve its yield is on increase. The plant is a perennial herb but is grown as an annual plant. It is a branching, herbaceous, hairy plant with weak trailing stems. The leaves are hairy and vary in size. It bears yellow flowers in clusters. The fruits are round to lobed and they vary in size and colour ranging from red, pink or yellow when ripe. The shape of the fruit differs per cultivar. Flat, slightly curved, hairy, light brown seeds are produced (Tigchelaar, 1978).

It is a succulent and fleshy vegetable which can reach a height of over two metres. In some regions, e.g. South America, however, the same plants can be harvested for several years in succession. The first harvest is possible 45–55 days after flowering, or 90–120 days after sowing (Tindall, 1968). Tomato is a warm-season crop that is sensitive to frost. An average daily mean of 20° to 24°C is optimum for growth, yield and fruit quality. Fruit set and quality are poor at temperatures below 12°C and 35°C (Simon and Sobulo, 1974). As it is a relatively short duration crop and gives a high yield, it is economically attractive and the area under cultivation is increasing daily (Varela *et al.*, 2003). Nigeria is the second largest producer of tomato in Africa second only to Egypt and 13th in the world, and produces 6 million tonnes of tomato annually prior to 1990 (Erinle, 1989). Tomato is cultivated almost all over Nigeria (Olanrewaju and Swamp, 1980). This work elucidates the mutagenic effects of the chemical mutagen (sodium azide) on three accessions of *Solanum lycopersicon* with the aim to ascertain the effects of this mutagen on the morphology and yield.

MATERIALS AND METHODS

Three accessions of tomato were used, accession (NGB01302) was obtained from National Centre for Genetic Resources and Biotechnology (NACGRAB) Ibadan, Nigeria. Two accessions (Tropimech and Tima) were obtained from National Horticultural Research Institute (NIHORT) Ibadan, Nigeria. The M1 seeds were then subjected to treatment with sodium azide.

Three concentrations of Sodium azide (1mM, 2mM and 4mM) were prepared in 1.0M phosphate buffer of pH 4. The seeds of each accession were soaked in the solutions of the mutagen separately for 4 hours at room temperature. The container

was agitated periodically to ensure even take up of mutagen by the seeds. After the holding period, the seeds were rinsed under tap water to remove excess mutagen. The seeds of both the treated and control were then sown in pots for twenty (20) weeks. Each treatment was replicated three times using completely randomized design (CRD) (Adamu *et. al.*, 2007).

DATA COLLECTION

Data were collected on the following growth and yield parameters: seedling germination, plant height, number of leaves per plant, number of branches, root length, number of flowers, number of fruit and fruit weight.

RESULTS AND DISCUSSION

RESULTS

Germination of seeds

Germination percentage at day 8 which is the estimate of variability of a population of seeds was 100% in the control; however, variations were observed in the treated. Tropimech had 40% in the treatment with 1mM, while 50% germination rate was observed in both 2mM and 4mM respectively. NGB01302 had an 85% germination rate in treatment with 1mM and 65% in 4mM. TIMA accession had 65% germination rate in 4mM and 55% in 2mM treatments respectively (Table 1).

MORPHOLOGICAL EFFECTS

Plant height

Measurements of some quantitative effects of the mutagen on the plants began at 4WAP and were terminated at 20WAP. At 4WAP, Tropimech control had the highest plant height (9.75cm) followed by NGB01303 at 4mM (8.07cm) while the lowest was observed in TIMA, 2mM (3.10cm). At 20WAP, the highest plant height was recorded in 1mM of the Tropimech accession (40.10cm) while the least was observed in TIMA at 2mM (8.93cm) (Table 1).

Number of leaves

At 4WAP, the highest number of leaves was observed in the control of Tropimech (12.00) followed by NGB01302 with 4mM and 1mM (9.50). The least number of leaves was recorded in TIMA at 4mM (4.00). The highest number of leaves at 20WAP was recorded in Tropimech (113.25) while the least was in TIMA (27.00) (Table 1).

Number of branches

The highest number of branches was recorded in Tropimech 1mM (8.75) followed by the treatment at 4mM (8.00) in the NGB01302 accession. The lowest was recorded in the 2mM treatment (3.00) of the TIMA accession. At 20WAP, Tropimech had the highest number of branches (17.50) while TIMA had the least number of branches (5.25) (Table 1).

Flower number

Tropimech (10.00) produced the highest number of flowers in treatment with 1mM in the study. Treatment with 2mM did not produce flower while the control produced 2.00. In NGB1302, 4mM had the highest number of flowers (3.50). The highest number of flowers in TIMA was recorded in the control (3.00) (Table 1).

Root length

The highest root length (10.38) was recorded in the Tropimech accession with 1mM treatment followed by the NGB01302 (7.53) while the least was obtained in TIMA (2.90) with treatment of 4mM at the point of termination. In the control, Tropimech had the highest (5.40) while NGB01302 had the least (Table 1).

Fruit number

The highest number of fruit was found in the control of Tropimech, treatment with 2mM did not bear fruit. The control of Tropimech, NGB01302 and Tima were the highest in terms of number of fruit (12.00), (8.00) and (6.00) respectively. Between the accessions treated with 1mM, Tropimech was the only one with fruit (3.5) while the other accessions (NGB01302 and Tima) did not bear fruit (Table 1).

Fruit weight

The highest fruit weight was recorded in the Control of Tropimech (12.86g) followed by Control of NGB01302, NGB01302 treated with 1mM Sodium azide was the least. It was impossible to measure the fruit weight of Tropimech (2mM), NGB01302 (1mM), Tima (1mM, 2mM, 4mM) as they did not produce fruit (Table 1).

Heritability (h^2_B) and Genetic Advance

The genotypic variance, phenotypic variance, heritability in broad sense, expected genotypic advance (GA) for all parameters studied are presented in table 5.

The plant height showed a low heritability of 9.68% and a low expected genetic advance of 3.48%.

The leaf number of the three accessions had a low heritability of 6.91% but exhibited high genetic advance of 21.91%.

Branch number and root length both had low heritability and exhibited low genetic advance as well.

Number of flower, number of fruit and fruit weight had moderate heritability of 55.86%, 37.50%, 56.44% respectively. The genetic advance estimated for the number of flowers and fruit weight was moderate while that of fruit number was low.

Table 1
Morphological effect of sodium azide on tomato (*Solanum lycopersicon*)

Variety	Treatment	Percentage germination	Seedling height (cm)	Root length	Number of leaves/plant	Height at maturity (cm)	Number of branches/plant	Number of flower/plant	Fruit number/plant	Fruit weight /plant(g)
Tropimech	0mM	100 ^a	9.75 ^a	5.40 ^b	67.00 ^{ab}	32.4 ^d	16.00 ^a	2.00 ^b	12.00 ^a	12.86 ^c
	1mM	40 ^b	4.76 ^b	10.38 ^a	113.25 ^a	40.10 ^a	17.50 ^a	10.00 ^a	3.50 ^b	8.73 ^b
	50 ^b	3.75 ^b	5.58 ^b	67.75 ^{ab}	22.46a	0	0	0	0	0
	4mM	50 ^b	3.73 ^b	3.10 ^b	54.00 ^b	23.40 ^a	9.00 ^c	4.00 ^b	1.50 ^{bc}	2.33 ^c
NGB01302	0mM	100 ^a	6.00 ^a	3.00 ^c	63.00 ^b	34.40 ^b	10.00 ^b	2.00 ^a	8.00 ^a	0
	1mM	85 ^a	6.66 ^a	7.53 ^a	88.75 ^b	38.54 ^b	10.50 ^b	0	0	1.01 ^b
	2mM	80 ^a	6.67 ^a	4.55 ^b	82.25 ^b	31.88 ^b	10.50 ^b	0.50 ^b	1.00 ^b	0.61 ^b
	4mM	65 ^a	8.07 ^a	2.98 ^c	148.50 ^a	53.06 ^a	15.25 ^a	3.50 ^a	1.00 ^b	9.85 ^a
Tina	0mM	100 ^a	6.25 ^a	4.45 ^b	44.00 ^a	20.30 ^a	8.00 ^a	3.00 ^a	6.00 ^a	7.45 ^a
	1mM	57.5 ^b	3.46 ^{ab}	8.00 ^a	43.00 ^a	20.51 ^a	7.75 ^a	0	0	0
	2mM	55 ^b	3.10 ^b	4.18 ^b	27.00 ^a	8.93 ^b	5.25 ^a	0	0	0
	4mM	65 ^a	4.22 ^{ab}	2.90 ^b	33.25 ^a	18.35 ^a	5.50 ^a	0	0	0

Means with the same letter along a column are not significantly different.

Table 2

Mean square of the effect of sodium azide on tomato (*Solanum lycopersicon*)

Source of variation	Df	Seedling height (cm)	Root length	Number of leaves/plant	Height at maturity (cm)	Number of branches/plant	Number of flower/plant	Fruit number/plant	Fruit weight /plant(g)
Replication	3	5.62 ^{ns}	6.17 [*]	47.70 [*]	243.11 ^{ns}	199.83 [*]	24.70 [*]	0.60 ^{ns}	1.74 ^{ns}
Variety	2	26.98 ^{***}	14.37 ^{***}	50.27 ^{***}	2024.56 ^{***}	311.27 ^{***}	241.33 ^{***}	42.33 ^{***}	78.01 ^{***}
Treatment	3	18.64 ^{***}	79.70 ^{***}	15.68 ^{***}	341.57 ^{***}	18.29 ^{***}	311.91 ^{***}	171.38 [*]	216.16 ^{**}
Treatment/variety	6	12.73 ^{***}	2.26 ^{**}	15.43 ^{***}	206.81 ^{***}	25.54 ^{***}	20.30 ^{**}	28.88 ^{**}	18.02 ^{**}
Error	35	2.90	0.54	7.54	80.96	17.31	2.23	0.711	2.23

*=Significant P<0.05; ns=non significant (p>0.05).

=significant (p<0.01), *=non significant (p>0.01).

Table 3

Mean value of effects of sodium azide on tomato (*Solanum lycopersicon*)

Variety	Seedling height (cm)	Root length	Number of leaves/plant	Height at maturity (cm)	Number of branches/plant	Number of flower/plant	Fruit number/plant	Fruit weight /plant(g)
Tropimech	5.49 ^{ab}	6.32 ^a	75.50 ^b	29.59 ^b	16.18 ^{at}	11.68 ^a	4.75 ^c	6.26 ^a
NGB01302	6.85 ^a	4.45 ^b	95.62 ^a	39.46 ^a	12.12 ^b	6.188 ^b	3.00 ^{ab}	3.76 ^{ab}
Tima	4.25 ^b	5.14 ^{ab}	36.81 ^c	17.02 ^c	7.37 ^c	4.188 ^b	1.50 ^b	0.87 ^b

Means with the same letter within the same column are not significantly different

Table 4

The effects of sodium azide on three varieties of tomato (*Solanum lycopersicon*)

Treatment	Seedling height (cm)	Root length	Number of leaves/plant	Height at maturity (cm)	Number of branches/plant	Number of flower/plant	Fruit number/plant	Fruit weight /plant(g)
0Mm	7.33 ^a	4.33 ^b	58.00 ^b	29.03 ^{ab}	12.66 ^a	14.00 ^a	4.30 ^b	10.05 ^a
1mM	4.96 ^b	8.99 ^a	81.00 ^a	33.05 ^a	13.16 ^a	8.66 ^a	8.90 ^a	3.66 ^b
2Mm	4.50 ^b	4.89 ^b	59.00 ^b	21.09 ^b	10.50 ^a	3.83 ^b	4.89 ^a	0.33 ^b
4Mm	5.34 ^b	3.03 ^b	78.00 ^a	31.60 ^a	11.25 ^a	2.91 ^a	3.03 ^a	1.25 ^b

Means with the same letter within the same column are not significantly different

Table 5

Estimate of heritability of the effect of sodium azide on three varieties of tomato

PARAMETER	σ^2_g	σ^2_p	h^2	GA
Plant height	303.83	3137.65	9.68	3.48
Number of leaves	1636.89	23679.44	6.91	21.91
Number of branches	4.16	610.07	0.68	0.35
Number of flower	100.96	179.00	55.86	15.40
Number of fruit	14.94	39.84	37.50	4.88
Fruit weight	101.15	179.20	56.44	15.56
Root length	0.89	26.09	3.41	0.35

 $H^2 = 0-30\%$ (Low)

30-60% (Moderate)

60 - above (High)

(Reddy *et al.*, 2001)

GA = <10% (Low)

10-20% (Moderate)

>20% (High)

(Johnson *et al.*)

DISCUSSION

Treatment with sodium azide exhibited a general reduction and lateness in the germination of seeds when compared with the control. The control germinated 4 days after planting while seeds treated with sodium azide germinated after day 5 and this was observed in all the accessions. In the Tropimech, there was 100% seed germination in the control, there was reduction in the seed germination in the treatments. Treatment with 1mM had 40% germination, 2mM had 50% germination and 50% germination in treatment with 4mM and this showed a great reduction in germination. NGB01302 also had a 100% germination in the control, 1mM (85%), 2mM (80%), 4mM (65%) which are also less than the control. More so in the third accession (Tima), the control had 100% germination, 1mM had 57.5% germination, 2mM (55%) and 4mM (65%) which are also lower than the control. Earlier work by Pande and Khetmalas (2012) reported that germination of seeds and survival of seedlings is strongly inhibited with increasing concentration and treatment duration. He explained that there was reduction in the germination and survival percentage with increasing concentrations for both chemicals in the first generation. Blixt (1970) opined that the inhibition in seedling growth might be due to the gross injury caused at cellular level either due to gene controlled biochemical processes or acute chromosomal aberrations or both. Evans and Sparrow (1961) suggested that the chromosomal damage and inhibition of cell division are the chief causes of reduced seedling growth. Khan and Goyal, 2009; Chowdhury and Tah (2011), explained that the decrease in seed germination induced by mutagenic treatments may be the result of damage of cell constituents at molecular level or altered enzyme activity. Micco *et al.*, (2011) have correlated seed germination with abnormalities in mitotic cycles and in metabolic pathways of the cells.

Decrease in plant height, number of leaves and number of branches were observed with increase in sodium azide concentration in the Tropimech and Tima accessions at the point of termination. This agrees with the work of Adamu and Aliyu (2007), Pande and Khetmalas (2012), Shagufta *et al.* (2013). Ignacimuthu and Babu (1988), Cherry *et al.* (1962) also reported that change in the specific activity of quite a few enzymes and physiological injury induced in the seeds and seedlings may be responsible for the decrease in height. Adamu and Aliyu (2007) showed that there was general decrease in germination percentage, seedling height, root length, number of leaves per seedling, seedling survival, height at maturity, number of fruit with increase in mutagenic concentration. The results obtained in NGB01302 in this study however differ from the above assertion. An increase in all the parameters was observed with increasing concentration of sodium azide at the point of termination. Krishna *et. al* (1984) opined that the greater sensitivity at higher concentration level was attributed to various factors such as changes in the metabolic activity of the cells, inhibitory effects of mutagens and disturbance of the balance between promoter and inhibitors of growth regulators.

The flower produced by Tropimech treated with 1mM sodium azide developed into fruit at week 15. However, flower senescence was observed in the treated accessions which consequently led to less production of fruit, this was in accordance to the work of Adamu *et. al* (2002) and Adamu and Aliyu (2007), the result of their work showed that all the traits studied were affected by sodium azide treatment. The decrease in seedling emergence, seedling height, root length, seedling survival, height at maturity and fruit yield per plant with increasing mutagen concentration has been reported by Aliyu *et al.*, (2002) in their mutagenesis studies on groundnut. In all growth parameters studied, Tropimech and Tima had a general decrease with increasing concentration of sodium azide. However, NGB01302 had an increase in all parameters studied with increasing concentration.

CONCLUSION

The treatment of the three tomato accessions with sodium azide exhibited a general increase in all parameters studied. M1 seeds showed heritable changes which can be used for selection of beneficial mutant leading to variability in tomato, the estimate of heritability has proved that these tomato accessions will perform well when selected. Low concentration of sodium azide can therefore be utilized to trigger traits that can help increase farmers' turnover which can ultimately lead to the improvement of tomato accessions.

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The book *Nanotechnologies in Food and Agriculture*, first edition, contains 14 chapters dealing with fundamental and applicative overviews on new and emerging nanotechnologies. Topics include nanoagrochemicals (3 chapters: Nano-fertilizers for Balanced Crop Nutrition; Nano-fertilizers and Their Smart Delivery System; Perspectives in Nanocomposites for the Slow and Controlled Release of Agrochemicals: Fertilizers and Pesticides), nanotechnologies for food processing, packaging, and storage, crop improvement and plant disease control (6 chapters: Strategic Role of Nanotechnology in Fertilizers: Potential and Limitations; Nanotechnology Applied in Agriculture: Controlled Release of Agrochemicals; Nanobiotechnology Strategies for Delivery of Antimicrobials in Agriculture and Food; Nano-developments for Food Packaging and Labeling Applications; Emerging Role of Nanocarriers in Delivery of Nitric Oxide for Sustainable Agriculture; Nanotechnology in Foods), Nanoecotoxicology (2 chapters) as well as individual chapters on nanobiosensors (Strategic Role of Nanobiosensor in Food: Benefits and Bottlenecks), plant genetic transformation (Nanoparticles-Based Delivery Systems in Plant Genetic Transformation) and waste treatment (Nano-enhanced Biological Treatment of Agricultural Wastewater). Each chapter contains very rich and useful information, including comprehensive references. Each chapter is very well written by well known scientist(s) in their fields. The book is not very rich in illustrations but the quality of the text partly overcomes this aspect. The book, useful and needed both by PhD students and already experienced scientists, should be found accessible in every (significant) faculty or research institute dealing with this still emerging field.

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