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Characterization of Cyanobacterial Strains Isolated from Soils Polluted with Insecticides

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ABSTRACT

This research aimed to isolate and characterize cyanobacteria that have the ability to grow in the polluted soil with insecticide and evaluation of their efficiency in fixing nitrogen and dry weight. Cyanobacteria were isolated from samples of soil site polluted with insecticides in Kafr El-Sheikh, Governorate. Cyanobacteria were purified by different purification methods. Twelve cyanobacterial isolates were identified according to standard methods based on cultural and morphological characters of cyanobacteria as following: color of culture, morphology of thallus, vegetative, reproductive cells and heterocyst. Heterocyst from isolates of cyanobacteria were observed in *Anabaena* sp., *Nostoc* sp., *Oscillatoria* sp. and *Chroococcus* sp. *Nostoc* sp. was of the highest frequency (66.6 %) among the isolated cyanobacteria. While, *Oscillatoria* sp. and *Chroococcus* sp. were less frequency (8.3%). Dry weight and nitrogen fixation of cyanobacteria were determined for efficiency evaluation of these isolates. The superior cyanobacterial isolates, which isolated from soil polluted with insecticides were identified by 16S rRNA as *Nostoc muscorum* and *Anabaena oryzae*.

Keywords: cyanobacteria, nitrogen fixation, insecticides .

INTRODUCTION

Recently, cyanobacteria have been discovered to be a main source of many active compounds such as extracellular products and secondary metabolites (Afify and Ashour 2018) such as vitamins, enzymes, carbohydrates, peptides and amino acids, which have been found to increase plant crops (Singh *et al.* 2016). Zulpa *et al.* (2003) and Sammauria *et al.* (2020) found that cyanobacteria are different communities of photosynthetic prokaryotes and abundant in soil, water, as well air ecosystems (Seckbach, 2007). Currently, the farmers extensively used insecticides for the protection of plants from insects (Parte *et al.* 2017). While, insecticides has enhanced in product yield and prevented insect-borne diseases (Verma *et al.* 2014). But there are, the problems associated with the use of these chemicals have also increased. It has become a major cause of environmental pollution such as soil and water (Rani and Dhania 2014). For cyanobacterial process and an adequate population biological N₂ fixation is very important. Cyanobacteria are group of photosynthetic prokaryote and gram negative bacteria (El-Saadny, 2013 ; El-Zawawy, 2016; Abou Elatta, 2018; Zaki, *et al.* 2021). In addition, insecticides effects on growth, nitrogen fixation and photosynthesis in cyanobacteria (Mohapatra *et al.* 2003; Jha and Mishra 2005; Prasad *et al.* 2005; Chen *et al.* 2007). Insecticides represent the greatest proportion of pesticides used in developing countries. Chlorpyrifos used since 1960s, for the control of crop from insects (Bicker *et al.*, 2005) but contaminated aquatic and terrestrial ecosystem and public health because it has long half-life and high residual concentration (Nawaz *et al.*, 2011). Chlorpyrifos is remains biologically active in soil for periods ranging from

twenty to eighty days and is moderately persistent, from ten to sixty days (Lakshmi *et al.* 2008). Residues of applied pesticides stay in the environment (air, soil, ground and surface water) for variable periods of time (Gavrilescu, 2005 ; Tariq *et al.*, 2007). This research aimed to isolate and identify cyanobacteria that have the ability to grow in the polluted soil with insecticides and evaluation of their efficiency in fixing nitrogen and dry weight.

MATERIALS AND METHODS

Source of soil samples :

Soil samples were collected from different locations at Kafr ElSheikh Governorate cultivated with rice and polluted with insecticides. The collected soil samples were used as a source of cyanobacterial isolates. Some chemicals and physicals analyses of soil (Piper 1950 and Jackson 1973) are previously presented in Abou Elatta *et al.*,(2023).

Isolation of cyanobacteria:

The method of preparation of the soil which were collected from rice fields polluted with the pesticides was carried out according to the Oxford Manual of Culture Media (Manual, 1990). BG11 medium (nitrogen – free medium) was used (Abdel-Razek *et al.*, 2019) the soil samples (little milligrams) were spread in the Petri dishes containing BG11 medium (Black *et al.*, 1965) and dishes incubated at 28-30°C with constant lighting of 2500 lux until appearance of cyanobacterial growth. After incubation cyanobacteria were purified by standard cyanobacterial purification techniques (Desikachary, 1959 & El-Ayouty and Ayyad, 1972). The purification techniques according to Watanabe liquid medium (Watanabe *et al.*, 1951) without nitrogen sources.

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Morphological identification of isolated cyanobacteria:

Cyanobacterial isolates were identified according to standard methods of Bergey's Manual of Systematic Bacteriology (2001) based on cultural and morphological characters of cyanobacteria, *i.e.* color of cultures, morphology of thallus, vegetative, reproductive cells and heterocyst.

Molecular identification of cyanobacterial isolates:

Only the most efficient cyanobacterial isolates were identified by molecular 16S rRNA genes of approximately 1500bp length in Agricultural Genetic Engineering Research Institute (AGERI) Agriculture Research Center, PO Box 12619, Giza, Egypt. RNA genes contain regions of variable DNA sequence that are unique to the species carrying the gene. The species identity of an unknown bacterium may therefore be deduced from its unique rRNA gene sequence.

Selection of the most efficient cyanobacterial isolates in fixing nitrogen and dry weight :

Selection of the most efficient cyanobacterial isolates in fixing nitrogen and dry weight. Each cyanobacterial isolate was cultivated separately for 21 days on suitable medium (modified Watanabe liquid medium) at 28-30°C with constant lighting of 2500 lux. The dry weight and fixed nitrogen were determined for each cultivated isolate. The most efficient isolates were selected.

Preparation of cyanobacterial inoculum:

Liquid cultures of cyanobacterial isolates were prepared using Modified Watanabe liquid medium with incubation at 28-30°C under continuous illumination (2500 lux).

Total nitrogen determination:

using the micro-kjeldahl method according to Jackson (1973), total nitrogen was determined for the cyanobacterial isolates .

RESULTS AND DISCUSSION**Distribution and morphology of cyanobacterial isolates:**

Twelve cyanobacterial isolates were obtained from a soil sample polluted with insecticides in Kafr El- Sheikh as described by El-Gamal *et al.* (2008). These twelve isolates were found to be belonging to four genera (*Anabaena*, *Nostoc*, *Oscillatoria* and *Chroococcus*). Data presented in Table (1) indicate that these genera varied in their densities

and frequency in the collected soil sample either in broth or solid medium (Staub, 1961). Among the twelve isolates eight isolates were found to be belonging to *Nostoc* Genus, representing 66.6% of the total isolates. Moreover, two isolates were identified as *Anabaena* genus among the twelve isolates representing 16.6% of the total isolates. Furthermore, each of *Oscillatoria* and *Chroococcus* genera was found to represent 8% of the total isolates. These results may indicate that *Nostoc* is the most dominant genus among the detected genera in the insecticides polluted soil. Similar results were obtained by Venkataraman (1981) & Roger and Ardales (1991).

Table 1. Densities of the isolated cyanobacterial genera in the collected soil sample polluted with insecticides

Origen soil samples	Cyanobacteria genera	No. of Isolates	Genera Frequency (%)
Kafr El-Sheikh Governorate, Egypt	<i>Anabaena</i> sp.	2	16.6
	<i>Nostoc</i> sp.	8	66.6
	<i>Oscillatoria</i> sp.	1	8.3
	<i>Chroococcus</i> sp.	1	8.3
Total number of isolates	4	12	100

The morphological characters of the twelve cyanobacterial isolates were studied based on the phenotypic properties, appearance and color of cultures in addition to the microscopic examination. The characteristics of the isolated cyanobacterial genera are presented in Table (2). In the Bergey's Manual of Systematic Bacteriology (2001) cyanobacteria contain five groups or subsections. According to the dichotomous key: the morphotype is unicellular or trichome; presence or absence of differentiated cells (Naz *et al.* 2004; Pinevich 2008; Shariatmadari & Riahi 2010; and Komárek *et al.* 2014). Individual cyanobacteria of all isolates were identified by cultural appearance (color, shape of colonial aggregates) in addition to its distinct of cells or filaments, heterocysts and akinetes.

Results in Figure (1) represent light micrographs of cyanobacterial isolates grown on nitrogen free medium (BG11) for 30 days.

Table 2. Cultural and microscopic characterization of cyanobacterial isolates

Cultural color	Thallus morphology	Vegetative Cell			Heterocysts			Akinetes			Cyanobacteria Identified Name	
		Shape	Width (µm)	Length (µm)	Site	Width (µm)	Length (µm)	Shape	Shape	Width (µm)		Length (µm)
Dark green	Filaments	oblong	3.2-4.8	4.1-4.5	Terminal and intercalary	3.2-4.8	4.1-4.5	Subspherical	oblong or oval,	4.1-6.3	5-6.3	<i>Nostoc paludosum</i>
Green	Filaments	Barrel	4-5	4-5	Terminant ercalary	3-3.5	4-4.5	Subspheric al conical	3-6 in series, sub-spherical	5-6	6-7	<i>Anabaena oryzae</i>
yellowish green later brownish	Filaments	quadrate to short barrel	2.7-3.6	2.7-4.1	Terminal	4.1-4.5	4.1-5	Spherical	-	3.6-7.2	4.1-6.5	<i>Nostoc entophyllum</i>
Blue-green, or brown; black	Filaments	Subglobose to barrel	2-3	3-4	Intercalary or terminal	5-6	5-7	Globose or ellipsoid	Spherical	10	10	<i>Nostoc pruniforme</i>
Oliveceous green, blue green	Filaments	Cylindrical	3.6-4.1	4.1-5	Terminal	4.2-5	4.2-5.4	Spherical	Ellipsoidal or oblong	5.0-6.3	5-8.1	<i>Anabaena gelatinicola</i>

Cont. Table 2. Cultural and microscopic characterization of cyanobacterial isolates

Cultural color	Thallus morphology	Vegetative Cell			Site	Heterocysts		Shape	Akinetes		Cyanobacteria Identified Name
		Shape	Width (µm)	Length (µm)		Width (µm)	Length (µm)		Width (µm)	Length (µm)	
Dark green	Filaments	Barrel, granular, yellowish	5-6	5.5-7	Terminal-Intercalary	-	-	-	-	-	<i>Nostoc muscorum</i>
Dark green	Filaments	apical cells sometimes slightly larger, oval	1.6-1.8	2.6-3	Terminal	4.6-6.2	5.2-8	Intercalary	6-8.5	6.5-11	<i>Nostoc viride</i>
Young cells palegreen, older ones brown	Filaments	quadratic, oblong or barrel	4.1-5.4	3.6-5.4	-	4.5-6.8	5-7.2	terminal or intercalary, cylindrical	-	-	<i>Nostoc Rivulare</i>
blue green	Slimy-gelatinous	Spherical	3-4	3-4	-	-	-	-	-	-	<i>Chroococcus minor</i>
Dark green	Solitary and straight	Solitary	4-8.2	2.5-3.1 4-7.1	-	-	-	-	-	-	<i>Oscillatoria brevis</i>

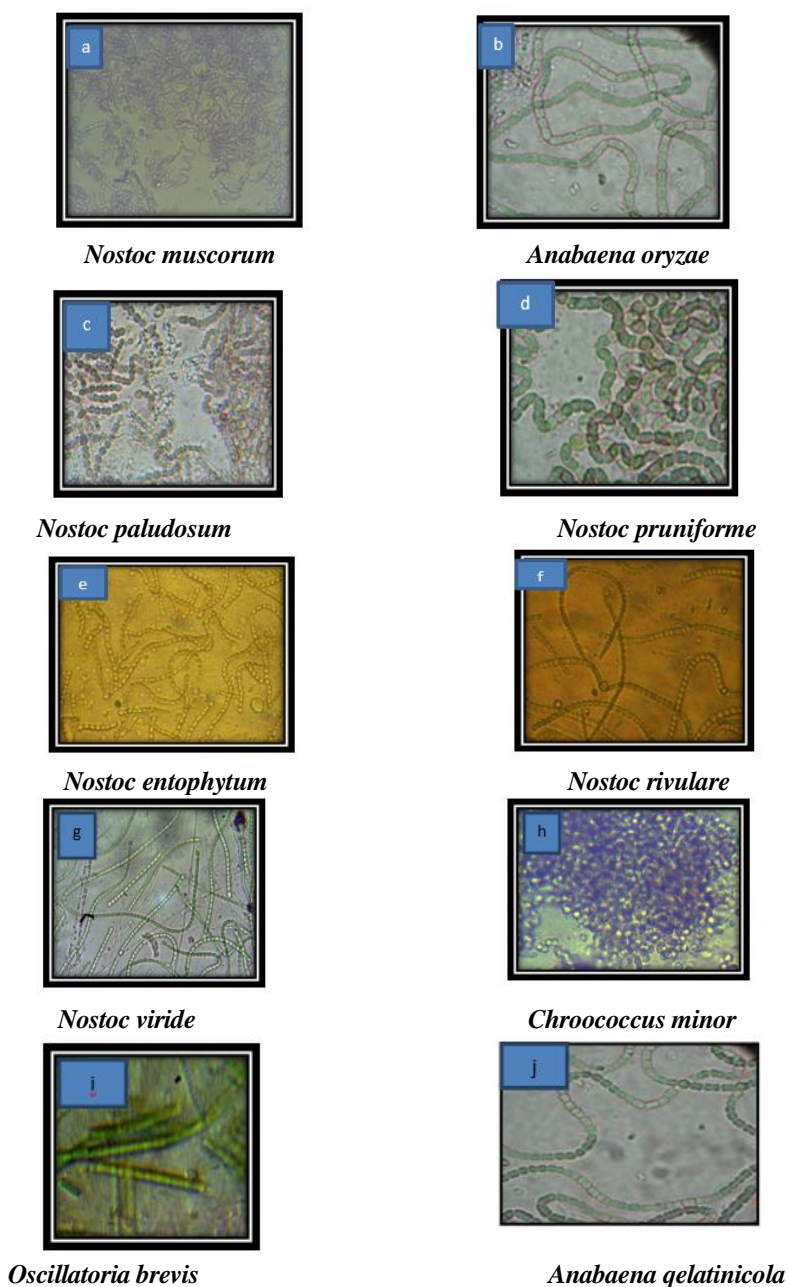


Fig. 1. Light micrographs of selected cyanobacterial strains.

Determination of dry weight of cyanobacterial isolates and their ability in fixing nitrogen:

The data presented in Table (3) indicate that the biomass dry weight of the cyanobacterial isolates and the fixed nitrogen in their liquid cultures increased gradually with increasing the incubation period. The highest values of dry weight and fixed nitrogen were recorded for all strains after inocubation for 21 days. The highest values of biomass dry weight (98 mg/100ml-culture) and fixed nitrogen (9.81 mg N/100 ml-culture) were recorded for *Anabaena oryzae* among all tested strains after incubation for 21 days. Whereas, the lowest value of biomass dry weight (60mg/100ml-culture) was recorded for *Nostoc verrucosum* and the lowest value of fixed nitrogen (4.27 mg N/100 ml-culture) was recorded for *Nostoc rivulare*. These results are in agreement with those obtained by Abou Elatta (2018) and Zaki et al. (2021) who found that the highest dry weight of cyanobacteria increased with increasing the incubation period.

Table 3. Dry weight (mg/100ml-culture) and nitrogen fixation (mg N/100 ml-culture) of the cyanobacteria strains.

cyanobacteria strains	Dry weight (mg/100ml-culture)		nitrogen fixation (mg N/100 ml-culture)			
	Incubation Period					
	7	14	21	7	14	21
<i>Oscillatoria brevis</i>	23	34	67	1.87	2.84	6.46
<i>Chroococcus minor</i>	26	51	65	1.91	2.83	4.18
<i>Nostoc paludosum</i>	31	57	79	2.33	4.14	9.55
<i>Anabaena oryzae</i>	52	74	98	3.39	6.94	9.81
<i>Nostoc pruniforme</i>	24	40	85	2.24	4.16	7.98
<i>Anabaena variabilis</i>	27	44	65	1.35	3.61	7.42
<i>Nostoc verrucosum</i>	28	40	60	2.12	4.58	5.81
<i>Anabaena qelatinicola</i>	30	45	76	2.46	3.89	4.34
<i>Nostoc entophytum</i>	42	55	62	3.11	5.45	8.06
<i>Nostoc rivulare</i>	26	46	66	1.41	2.26	4.27
<i>Nostoc viride</i>	33	62	72	2.41	3.35	4.35
<i>Nostoc muscorum</i>	48	65	91	3.25	5.35	9.81

Molecular identification of the most efficient cyanobacterial isolates:

Based on the results, it can be inferred that the most efficient isolates for nitrogen fixation ability and biomass production belonged to two strains of *Nostoc muscorum* and *Anabaena oryzae*. Molecular identification revealed that these two isolates contained 16S ribosomal RNA (rRNA) genes that were approximately 1500bp in length. The sequences were analyzed using BLAST program (<http://www.ncbi.nlm.nih.gov/BLAST>) and aligned using Align Sequences Nucleotide BLAST (Figure 2). These genes consist of regions of variable DNA sequence that are unique to the species harboring them. Therefore, the identity of an unknown bacterium can be determined from its distinct rRNA gene sequence. To do this, rRNA genes are first amplified using PCR technology. Subsequently, PCR cycle sequencing is conducted, and the rRNA sequence is determined with a capillary sequence analyzer. The resulting sequence is then compared to known rRNA sequences in Gen-Bank® and subjected to a rigorous review process for validation (as shown in Figures 3 and 4).

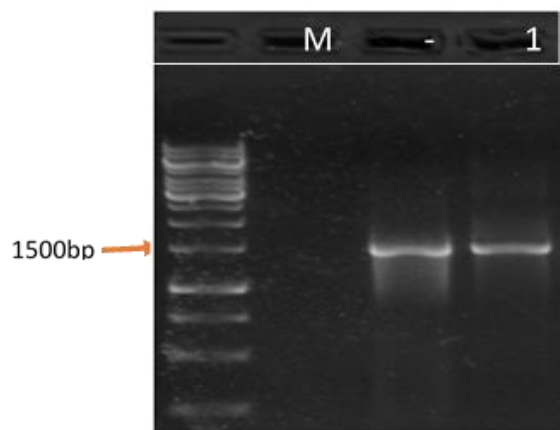


Fig. 2. The sequences nucleotide BLAST program.

Nostoc muscorum

GGAATTGCGATTGCTTACTATGCAGTCGAACGGGCTTTCGGTGATCTGGCGGCTCAGGATGAACGC
 TGGCGGTATGCTTAACACATGCAAGTCGAACGGTCTCTTCGGAGATAGTGGCGGACGGGTGAGTAAACGG
 TGAGAATCTAGCTTCAGGTCGGGGACAACCCTGGAAACGGTGCTAATACCGGATGTGCCGAAAGGTG
 AAAGATTATTGCTGAAGATGAGCTCGCTCTGATTAGCTAGTTGGTGTGGTAAAGAGCGCACCAAGGCG
 ACGTACAGTAGCTGGTCTGAGAGGATGATCAGCCACACTGGGACTGAGACACGGCCAGACTCCTACGG
 GAGGCAGCAGTGGGAATTTCCGCAATGGCGAAAGCCTGACGAGCAATACCGCGTGAGGAGGAAG
 GCTCTTGGTTGTAACCTCTTTCTCAGGGAATAAAAAAATGAAGGTACCTGAGGAATAAGCATCGGCT
 AACTCGTGCCAGCAGCCGCGTAATACGGAGGATGCAAGCGTTATCCGGAATGATTGGCGTAAAGCG
 TCCGCAGGTGGCACTGTAAGTCTGCTGTTAAAGAGCAAGGCTCAACCTTGTAAAGGCAGTGGAACTACA
 GAGCTAGAGTACGTTCCGGGCAGAGGGAATTCCTGGTGTAGCGGTGAAATGCGTAGAGATCAGGAAGAA
 CACCGGTGGCGAAAGCGCTCTGCTAGGCCGTAAGTACTGACTGAGGGACGAAAGCTAGGGGAGCGAATGG
 GATTAGATACCCAGTAGTCTAGCCGTAACGATGGATACTAGCGGTGGCTTGTATCGACCCGAGCCGT
 GCCGGAGCCAACGCGTTAAGTATCCCGCTGGGAGTACGCACGCAAGTGTGAAACTCAAAGGAATTGA
 CGGGGGCCGACAAAGCGGTGGAGTATGTGTTAATTTCGATGCCAACGCGAAAGAACCTTACCAAGACTTG
 ACATGTCGCAATCTTCTTGAAGGGAAGAGTGCCCTTAGGGAGCGCGAACACAGGTGGTGCATGGCTGTC
 GTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCGCTACG

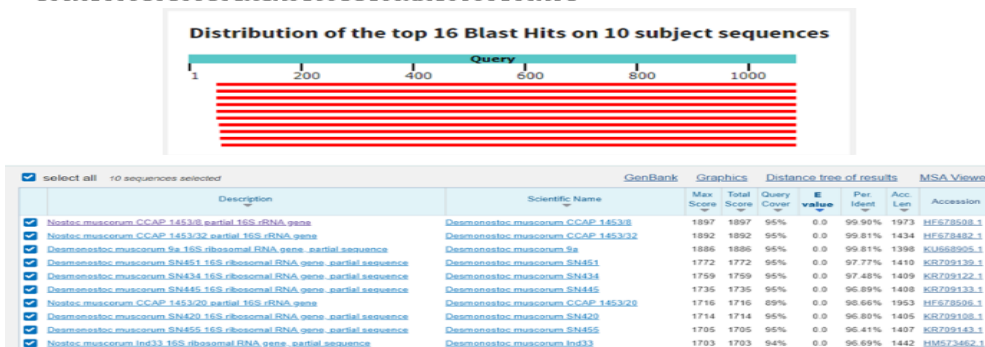


Fig. 3. Genetic bioformations background for the identification of *Nostoc muscorum*.

Anabaena oryzae

GAGATCGTGGCGGACGGGTGAGTAAACCGTGAGAATCTAGCTTCAGGTCGGGGACAACCCTGGA AACGGTGGCTAATACCGGATGTGCCGAAAGGTGAAAGATTTATTGCCTGAAGATGAGCTCGCTCTGATT AGCTAGTTGGTGTGGTAAGAGCGCACCAAGGCGACGATCAGTAGCTGGTCTGAGAGGATGATCAGCCAC ACTGGGACTGAGACACGGCCAGACTCTACGGGAGGCGAGTGGGGAATTTCCGCAATGGGCGAAA GCCTGACGGAGCAATACCGCGTGAGGGAGGAAGGCTCTTGGGTTGTAACCTCTTTCTCAGGGAATAAA AAAATGAAGGTACCTGAGGAATAAGCATCGGCTAACTCCGTGCCAGCAGCCGCGGTAATACGGGAGTGC CAAGCGTTATCCGGAATGATTGGGCGTAAAGCGTCCGAGGTGGCACTGTAAGTCTGCTGTTAAAGAGCA AGGCTCAACCTTGTAAAGGCAGTGGAAACTACAGAGCTAGAGTACGTTCCGGGACAGAGGGAATTCCTGG TGTAGCGGTGAAATGCGTAGAGATCAGGAAGAACACCGGTGGCGAAGCGCTCTGCTAGCCGTAACCTG AACTGAGGGGACGAAAGCTAGGGGAGCGAATGGGATTAGATACCCAGTAGTCTAGCCGTAACCGATG GATACTAGGCGTGGCTTGTATCGACCCGAGCCGTGCCGGAGCCAACCGGTTAAGTATCCCGCCTGGGGAG TACGCACGCAAGTGTGAAACTCAAAGGAATTGACGGGGGCCGCACAAGCGGTGGAGTATGTGGTTAA TTCGATGC.AACGCGAAGAACCTTACCAAGACTTGACATGTCGCGAATCTTCTTAAAGGGAAGAGTGCCT TAGGGAGCGCAACACAGGTGGTGCATGGCTGTCGTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCG CAACGAGCGCAACCCTCGTTTTTGTGGCAGCATTAAAGTTGGGCACTTAGAGAGACTGCCGGTGACAA ACCGGAGGAAGTGGGATGACGCTCAAGTCAGCATGCCCTTACGTCCTGGGTACACACGTAACAAT GCTACGGACAGAGGGCAGCAAGCTAGTGATAG



Fig. 4. Genetic bioformations background for the identification of *Anabaena oryzae*.

CONCLUSION

Generally, on the basis of the obtained results it can be concluded that cyanobacteria were found in soil polluted with insecticides. The detected cyanobacterial genera in such soil were *Anabaena* sp., *Nostoc* sp., *Oscillatoria* sp. and *Chroococcus* sp. *Nostoc* genus was found to be the most dominant genus among the detected genera in the insecticides polluted soil. Moreover, in liquid cultures of cyanobacteria the biomass dry weight and the fixed nitrogen increased with increasing the incubation period.

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خصائص سلالات السيانوبكتيريا المعزولة من عينات تربيته ملوثة بمبيدات الآفات الحشرية

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الملخص

تهدف هذه الدراسة إلى عزل السيانوبكتيريا من عينات تربيته ملوثة بمبيدات حشرية من أراضي مزروعة بالأرز في محافظة كفر الشيخ وقد تم الحصول على إثني عشرة عزلة نقيه. تم تنقية عزلات السيانوبكتيريا بالطرق المختلفة وتم تعريف عزلات السيانوبكتيريا طبقاً لطرق التعريف القياسية المزروعة (اللون) وكذلك المورفولوجية (شكل ولون الثالوس وحجم الهيتروسست بالإضافة إلى الخلايا الخضريه و التكاثرية) وإتضح عند تنمية أجناس السيانوبكتيريا أن كل من أنابينا ونوستوك وأوسيلاتوريا وكذلك كرووكوكس لها القدرة على تكوين الهيتروسست. وقد سجل جنس النوستوك أعلى نسبة في الانتشار بينما سجل جنس الأوسيلاتوريا والكرووكوكس أقل انتشاراً وذلك عند العزل من الأراضي الملوثة بهذه المبيدات الحشرية. وقد أثبتت جميع العزلات كفاءة في الوزن الجاف وكمية النترجين المثبت وكان أفضلهما سلالتى النوستوك والأنابينا ومن خلال التعريف الجزيئي ثبت أنهما يمثلان بنسبة ٩٩ و ٩٠% لكل من نوستوك مسكورم وأنابينا أوريذا. وبهذه النتائج يمكن أن نوصى باستخدام تلك السلالات من السيانوبكتيريا في التسميد الحيوى وخصوصاً في الأراضي الملوثة بمبيدات الآفات.