PRODUCTION OF ANTIMICROBIAL AGENTS FROM SOME BACILLUS ISOLATES

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ABSTRACT

A survey was undertaken for bacterial isolates capable of producing antimicrobial agents from different Egyptian soils. These isolates were tested against various species of microorganisms (G+, G-, spore formers, yeast and fungi) using inhibition zone determination technique and standard disks of antibiotics as reference.

One or more of the pathogenic bacterial strains used as tested organisms were affected by antimicrobial substances produced by bacterial isolates as indicated by inhibition zones formation. The effect of antimicrobial substances produced by bacterial isolates were compared with the effect of 11 antibiotic disks as reference. The most effective isolates against the tested organisms were identified by API 50 CHB system which indicate that these isolates were *Bacillus licheniformis*, *Bacillus amyloliquefaciens*, and *Bacillus subtilis*.

Whole sediment produced by the five bacterial strains were found to be safe when using as ration for albino rats.

Key words: Antimicrobial-Isolation and Identification - Biological assay experiment – *Bacillus*.

INTRODUCTION

A few decades after the introduction of antibiotics into clinical practice, resistance by pathogenic bacteria has become a major health concern. Indeed, while in the mid1970s infectious diseases were considered virtually conquered (**Breithaupt, 1999**), many Gram positive and Gram negative bacteria opportunistic pathogens were becoming resistant to virtually every clinically available drug (**Greenberg, 2003**).

In recent years, biological control of plant pathogens is considered as a potential control strategy as chemical control became undesirable. At present, effective management of plant diseases and microbial contamination in several agricultural commodities is generally achieved using synthetic pesticides. However, the continual and indiscriminate application of these chemical fungicides has caused health hazards in animals and human due to residual toxicity (Alemu, 2014).

Quite a large number of microorganisms have the capacity of producing bio-agents that can be used as antibiotics. From *Streptomyces*, **Ceylan** *et al.*, (2008) evaluated their inhibitory activities on seven microorganisms including multiple antibiotic resistant *Staph. aureus* and *Stenotrophomonas maltophilia*.

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Ogunmwonyi (2010) isolated ten putative *Streptomyces* strains and assessed them for antibiotic production and activity against a wide range of bacteria. Also, **Aparanji** *et al.*, (2013) collected six isolates of Actinomycetes from marine sediments in India. Among these six isolates, only one showed potent activity against all the bacteria and fungi tested.

Kawaguchi *et al.*, (2013) reported that among 151fungal isolates, 26 strains were found to reproduce anti- *C. albicans* activity in liquid medium.

Liasi *et al.*, (2009) found that three isolates of lactic acid bacteria (LAB) from the fermented food product identified as genus *Lactobacillus* (*L. casei* LA17, *L. plantarum* LA22 and *L. paracasei* LA02), the antibacterial agent produced by the isolates inhibited the growth of a range of gram-positive and gram-negative microorganisms.

Sixteen soil samples were collected by (**Kuta** *et al.*, **2009**) analyzed some soil samples for the presence of *Bacillus spp*. Antibiotic production tests indicated that nine *Bacillus* species out of twelve isolated could be used to produce antibiotics that had effect on the test organisms. However, *Paenibacillus polymyxa*, *B. sphaericus* and *B. laterosporous* had little or no effect on the tested organisms.

Sarker *et al.*, (2010) found that the gram-positive bacterium *B. subtilis* produces a large number of antibiotics, which are classified as ribosomal or nonribosomal. **Moshafi** *et al.*, (2011) screened a variety of microorganisms from environmental soil samples for antibiotics production. Bacterial isolates were found to have antibacterial and antifungal activity on the standard indicator species. **Tabbene** *et al.*, (2011) found that *B. subtilis* B38, isolated from soil showed antimicrobial activity against human pathogenic *C. albicans* species. Specific PCR primers revealed the presence of the bam C gene, which is involved in the biosynthesis of bacillomycin D. **Malnicheva** *et al.*, (2012) isolated bacterial strains with a high level of antimicrobial activity from soil and identified as *B. megaterium*. In submerged cultures, nine out of ten *B. megaterium* strains were found to produce antibacterial antibiotics differing in their spectra of action.

The aim of this investigation is an attempt to find antimicrobial principals naturally produced by some well-known bacteria which can replace the use of chemotherapeutic products.

MATERIALS & METHODS

Collection and preparation of soil samples:

In systematic screening program for the isolation of some target bacteria, 30 soil samples were collected from different locations from upper layer where most of the microbial activity usually takes place and where most of the bacterial population is concentrated.

Isolation of antimicrobial agents producing bacteria:

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Seventy isolates were obtained from soil samples by preparing serial dilutions from 10-1 to 10-10. The isolation of bacteria was made to whom which gave a clear zone of inhibition around their growth after 24 to 48 hours incubation period. The selection of the isolates was made only for those having a very wide zone of inhibition and those which never lost their potency when tested again in replicated inoculation , and those which were active , principally , against *Bacillus* spp.

Test microorganisms:

Sum of eleven different kinds of identified microorganisms were used in this study. Two G- bacteria (*E. coli* ATCC 25922 and *Salmonella typhimurium*, four G+ bacteria (*Listeria monocytogenes* ATCC 15313, *B. cereus* ATCC 13753, *Micrococcus luteus*, and *Staphylococcus aureus* ATCC 8095), Human pathogenic yeast *Candida albicans*, and fungi *Penicilium roquefortii*, *Aspeagillus niger*, *Fusarium semitectum*, and *Alternaria alternata* were used as test microorganisms for the detection of the antimicrobial activity. These strains were chosen because they have important sanitary position for human, animals and plants.

Antibiotics:

Eleven well known antibiotics were used *in vitro* diagnosis as reference to be compared in their action on different tested organisms with that of the isolated bacteria from soils. Claforan, Nebcin, Garamycin, Unacin, Tarivid, Rociphin, Fucidin, Cefobid, Lincocin, Velosef and Keflex. These eleven antibiotics were chosen because they were already used in medicinal treatments against a wide variety of diseases whether for human, animals, and plants. Nearly all had broadcast effects.

Antimicrobial activity assay

A- Disc diffusion assay for bacteria :

Luria-Bertani agar plates were inoculated with 1% tested organism, left to dry then antibiotic containing discs were applied. The plates were incubated for 24 hrs at the optimum temperature for each bacterial strain. The antibacterial activities of antibiotics were assessed by measuring the zone of inhibition (mm) against the tested bacteria (**Kumar** *et al* **2012**).

B- Disc diffusion assay for fungi :

Potato dextrose agar medium was used for determination of the antifungal effect of *Bacillus* isolates. The pH of the medium was adjusted to 5.6 ± 0.2 then autoclaved at 121°C for 15 min. Then constant number of spores of different fungi were added to 20ml of medium and poured in Petri dishes and let for drying and then the disks of *Bacillus* isolates containing known amount of supernatant (0.1ml) were placed on the surface of agar and incubated for appropriate time . Clear zone around each disc were examined after two days of incubation (**Farhana** *et al.*, **2011**).

Identification of *Bacillus* isolates

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The carbohydrate fermentation patterns of the selected five Bacillus isolates were determined with API CHB50 fermentation strips in CHB medium as specified by the manufactures instruction. For API kit method, sufficient growth of each isolate being identified was inoculated into the suspension medium supplied with the API Bacillus kit so as to produce an inoculum density equivalent to a MacFarland 2 standard. Separate suspension for use with the API API 50 CHB, was prepared at a density equivalent to a MacFarland 2 standard. Using sterile Pasteur pipettes, approximately 1 ml of the organism suspension was inoculated into each microtubule of the incubation tray. The API systems were inoculated in accordance with manufacturer's instructions. After inoculation, identification system was incubated at 30°C for 24-48 hrs and read after 24 hrs incubation, with a final reading made after 48 hrs incubation. All results were recorded on the work sheets provided and interpreted using the API Web database systems. To identify an organism, the API web software compares the profiles obtained with the profiles of taxa (Aruwa and Olatope, 2015).

The 16S rRNA gene sequence has been determined for a large number of strains. Gen Bank, the largest databank of nucleotide sequences, has over 20 million deposited sequences, of which over 90,000 are of 16S rRNA gene. This means that there are many previously deposited sequences against which to compare the sequence of an unknown strain.

Lastly, the 16S rRNA gene is universal in bacteria, and so relationships can be measured among all bacteria. In general, the comparison of the 16S rRNA gene sequences allows differentiation between organisms at the genus level across all major phyla of bacteria, in addition to classifying strains at multiple levels, including what we now call the species and subspecies level. The occasional exceptions to the usefulness of 16S rRNA gene sequencing usually relate to more than one well-known species having the same or very similar sequences (**Jill**, **2004**).

Media:

Luria Bertani agar medium (Atlas and Parks 1997)

Luria bertani agar medium was used to determine the antimicrobial effect of *Bacillus* isolates supernatant by disk diffusion method containing (gm/l);10.0 g Tryptone, 10.0 g NaCl,

5.0 g Yeast extract, 15.0 g Agar at pH 7.0 ± 0.2 .

Potato dextrose agar medium (PDA), Atlas and Parks (1997) containing (gm/l); 200.0 g Potato infusion, 20.0g Dextrose and 15.0 g Agar at pH 4.5 was used for fungal growth and test.

For feeding rats, a basal diet containing; Casein 15%, Mineral salts 4%, Vitamins 1%, Corn oil 7%, Sawdust 5% and Starch 68%. **Feeding experiment (biological assay):**

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Thirty weanling albino rats with average weight of 110 to 149 gm, obtained from Agriculture Research Center were acclimated to animal house condition housed in metabolic cages ($30 \times 40 \times 35$ cm), for one week prior the experiment.

The rats were housed in groups of five each in universal polypropylene cages at room temperature. Cages were kept in well aerated place with continuous artificial aeration all over the experimental period. Albino rats were fed on normal basal diet incorporated with certain amounts of the sediment obtained from growing different *Bacillus* strains on the basal medium.

Liver functions

The concentration of serum glutamic pyruvic (GPT) and glutamic oxaloacetic (GOT) transaminase were calourimetrically determined using transaminase kite according to the method of **Reitman and Frankel** (1957).

Kidney functions

Creatinine, bilirubin and urea were determined according to the methods described by (Gornall *et al.*, 1948; Doumas *et al.*, 1971; and Fawcett and Soctt, 1960) respectively.

RESULTS AND DISCUSSION

Isolates antagonism effect against the six tested bacteria and five tested fungi

Out of the 50 *Bacillus* isolates, the more effective five isolates, A6, A9, A13, A14 and A16 greatly inhibited *E. coli*, *L. monocytogenes*, *B. cereus*, *S. typhimurium* and *Staph. aureus*, respectively as shown in (table 1) were selected for further study.

When those five selected bacterial isolates were tested against some fungi and yeast they showed strong effect (Table 2).

Antibiotics susceptibility of the selected five Bacillus isolates

The five *Bacillus* strains were tested for their *in vitro* antibiotic susceptibility using disks of 11different reference antibiotics. The results obtained (**Table 3**) show the following: (i) all strains have showed a variable antibiotic resistance, (ii) all strains were resistant to Claforan, Cefobid, Fucidin. (iii) all strains were affected by Tarivid, Garamycin, and Nebcin (*L.monocytogenes* was not affected by Garamycin).

 Table (1): The more inhibititive effect of potent isolates against different pathogenic bacteria

Isolate	e Inhibition zone diameter (mm)					
No. B. M. Staph.				E	S	L
	cereus	luteus	aureus	.coli	.typhimurium	monocytogenes
A6	20	28	29	25	20	20
A9	28	30	20	35	33	25
A13	28	31	30	40	40	30
A14	25	30	30	35	38	32

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A16	20	26	30	45	28	20

- The diameter of the saturated disk was 9 mm - The medium used was nutrient broth. - Incubation period was 24hrs at 30°C.

	Inhibition zone diameter (mm):					
Isolate	<i>P</i> . <i>A</i> . <i>F</i> .				С.	
No.	roquefortii	n iger	Semitectum	alternata	albicans	
A6	44	29	44	55	60	
A9	31	25	56	35	38	
A13	44	29	64	30	44	
A14	54	44	42	40	45	
A16	34	29	44	23	38	

- The diameter of the saturated disk was 9 mm. - The medium used was nutrient broth.

- Incubation period was 24hrs at 30°C.

Table (3): Comparison between antibacterial effects of some tested
antibiotics and the *Bacillus* supernatants on the five tested
bacteria:

Antibiotics		Inhibition zone diameter (mm) : (antibiotics)					
disks	(µg/disk)	<i>B</i> .	Staph	L.	<i>S</i> .	<i>E</i> .	
		cereus	.aureus	monocytogenes	typhimurium	coli	
Keflex	30	N.E.	15	6	25	N.E.	
Velosef	30	9	N.E.	N.E.	9	N.E.	
Lincocin	30	18	14	13	25	N.E.	
Cefobid	75	N.E.	N.E.	N.E.	18	N.E.	
Fucidin	10	N.E.	N.E.	N.E.	N.E.	N.E.	
Rociphin	30	N.E.	14	13	30	21	
Unacin	30	18	16	15	18	N.E.	
Tarivid	10	30	26	33	40	45	
Gramycin	10	25	15	N.E.	23	31	
Nebcin	10	25	14	19	20	25	
Claforan	30	N.E.	N.E.	9	12	N.E.	
IsolateNo.	Dose	Inhibition zone diameter (mm)					
	(mg/Disk)	(Bacillus isolates supernatants)					
A.6	1	20	29	20	20	25	
A.9	2	28	20	25	33	35	
A.13	1	28	30	30	40	40	
A.14	1	25	30	32	38	35	
A.16	2	20	30	20	28	45	

- The diameter of the saturated disk was 9 mm. - The diameter of the standard disk was 6mm. - The medium used is nutrient broth. N.E. = no effect

Note: all antibiotics were added at the recommended dose usually implicated .

Identification of the five bacterial isolates using API 50 CHB kits:

Identification of the efficient selected five isolates to species level was carried out on the basis of their carbohydrates fermentation patterns obtained by API 50 CHB kits. According to the API database correlation these isolates were identified as two isolates (97.5%) *B. subtilis*, two isolates (99.9%) *B. amyloliquefaciens*, and one isolate (98.2%) *B. licheniforms*.

Genotypic characterization of the selected two Bacillus isolates:

The 16SrRNA gene sequences of isolates A6, and A13 were compared with sequences in the Gene-Bank database. A phylogenetic tree based on 16SrRNA sequences (**Fig. 1**) revealed affiliation of isolate A6 to the genus *Bacillus*, comprising 98 % identity to *B. sonorensis*, which is most closely related to *B. licheniformis*. While revealed affiliation of isolate A13 to the genus *Bacillus*, comprising 99% identity to *B. methylotrophicus*; Comparative genomic analysis showed only minor differences between this strain and the genome of the *B. amyloliquefaciens* subsp.

The effect of feeding rats on the whole sediment produced by tested *Bacillus* strains:

At the end of the feed experiment all groups except group 6 showed a slight decrease in weight, while the same group showed a slight increase in weight (Table 4). There are no significant differences between groups in the measurement of urea and enzyme alanine aminotransferase (ALT). While some differences were found between the groups in measuring creatinine (in group 4 decreased creatinine value while in group 5 it increased but other groups showed no significant differences). Enzyme aspartate aminotransferase (in group 2 was decreased while increased in group 4, 5 but other groups showed no significant differences (Table 5).

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As to blood picture, five groups showed an increase in hemoglobin and the number of red blood cells higher than the control. While some groups recorded a decrease in white blood cells (group 3, 4, 5, 6) and the other group recorded an increase in the number (group 2) compared with control group. (Table 6)

Group	Initial	Final	Total	Daily
-	weight (gm)	weight(gm)	gain(gm)	gain(gm)
1	126.8000	107.2000 ^b	19.6000 ^b	0.5540 ^b
	± 4.575	± 1.827	± 4.444	±0.126
2	126.8000	107.4000 ^b	19.4000 ^b	0.5500 ^b
	±4.363	±4.237	± 2.249	±0.063
3	127.4000	109.0000 ^b	18.4000 ^b	0.5220 ^b
	± 4.456	± 4.461	±3.155	±0.091
4	127.0000	108.0000 ^b	19.0000 ^b	0.5400 ^b
	± 5.966	± 6.244	± 3.633	±0.103
5	127.4000	112.0000 ^b	15.4000 ^b	0.4360 ^b
	±6.152	± 6.670	±2.731	± 0.076
6	127.2000	140.0000 ^a	12.8000 ^a	0.3620 ^a
	± 4.442	±9.612	± 5.228	±0.147

Table (4): The effect of feeding rats on the whole sediment produced by
tested Bacillus strains on body weights of female albino rats

1= control group

2 = group treated with isolate A6 3 = group treated with isolate A9

4= group treated with isolate A13 5= group treated with isolate A14 6= group treated with isolate A16

Table (5): The effect of feeding rats on the whole sediment produced by
tested Bacillus strains on liver and kidney functions of female
albino rats

	Urea	Creatinine	ALT(GPT)	AST(GOT)
Group	(mg/dl)	(mg/dl)	(U/L)	(U/L)
1	42.0000±3.60	0.9167 ^{ab} ±0.04	22.6667±1.20	51.3333 ^{ab} ±0.87
2	51.3333±4.05	0.9500 ^{ab} ±0.07	21.0000±0.57	41.3333 ^b ±3.37
3	47.3333±3.75	$0.8167^{ab} \pm 0.05$	23.0000±4.61	53.000 ^{ab} ±10.78
4	47.6667±3.37	0.7833 ^b ±0.04	21.3333±3.47	75.0000 ^a ±2.51
5	50.3333±5.92	1.0000 ^a ±0.02	23.3333±0.32	65.6667 ^a ±9.05
6	47.6667±2.72	0.933 ^{ab} ±0.03	25.3333±2.96	52.3333 ^{ab} ±9.38

2 = group treated with isolate A6 3 = group treated with isolate A9

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4= group treated with isolate A13 5= group treated with isolate A14 6= group treated with isolate A16

t	tested <i>Bacillus</i> strains on blood picture of female albino rats						
Group	Hb	R.B.C.	W.B.C.				
	(gm/dl)	(1000000/cm3)	(1000/cm3)				
1	10.900 d ±0.15	3.953 b ±0.10	6.3500 b ±633.11				
2	12.133 a ±0.23	4.556 a ±0.09	6.7500 b ±202.07				
3	10.933 d ±0.32	4.490 a ±0.21	5.1167 cd ±164.14				
4	11.833 abc ±0.21	4.423 a ±0.07	4.3000 d ±189.29				
5	11.266 bcd ±0.23	4.236 ab ±0.08	4.6833 d ±196.92				

4.170 ab ±0.13

Table (6): The effect of feeding rats on the whole sediment produced by tested *Bacillus* strains on blood picture of female albino rats

1= control group

6

3= group treated with isolate A9

2 = group treated with isolate A6 4= group treated with isolate A13

6.0000 bc ±407.22

5 = group treated with isolate AV 5 = group treated with isolate A14

 $11.000 \text{ cd} \pm 0.40$

6= group treated with isolate A16

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انتاج بعض المواد المضادة للميكروبات من بعض عزلات الباسيلس الخلاصة

أجرى حصر لبعض العزلات البكتيرية المنتجة للمضادات الحيوية من عينات مختلفة من الأراضى المصرية وتم اختبار تأثير هذه العزلات البكتيرية على مجموعة من السلالات البكتيرية السالبة والموجبة لجرام الممرضة للإنسان وكذلك بعض الفطريات والخمائر وذلك بتقدير المجال الرائق حول القرص المشبع بالمستخلص البكتيرى الخالى من الخلايا.

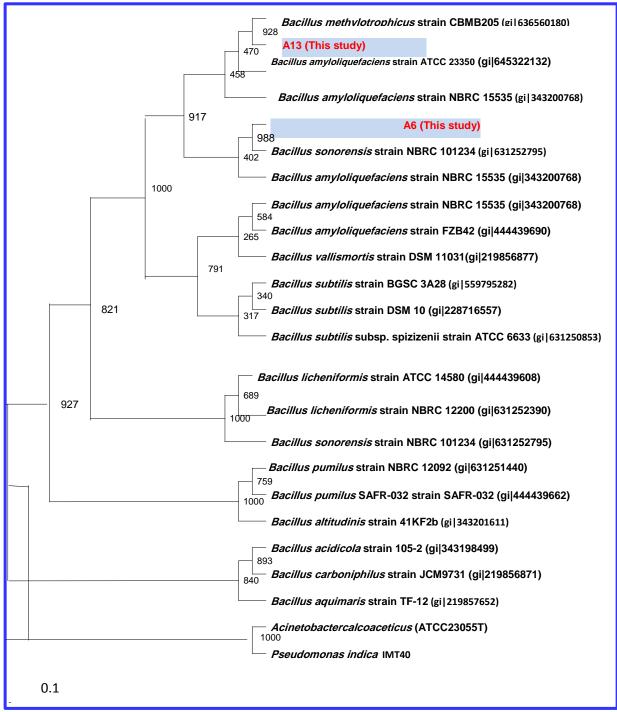
تأثرت واحدة أو أكثر من السلالات البكتيرية الممرضة والفطريات المختبرة بالمستخلص البكتيرى الخالى من الخلايا ثم تم اختيار أفضل خمس عزلات بكتيرية كان لها نشاط مضاد للبكتريا والفطريات.

تم مقارنة تأثير المواد المضادة المفرزة من العزلات البكتيرية مع ١١ مضاد حيوى وتم تعريف أقوى العزلات البكتيرية بنظام API 50 CHB والذى أوضح أن هذه العزلات هى Bacillus و Bacillus subtilis و Bacillus amyloliquefaciens و licheniformis كذلك تم تعريف أقوى عزلتين لهما تأثير مضاد للمبكر وبات بنظام 16sr RNA والذى اوضحت نتائجه

أن العزلات هي B. methylotrophicus & B. sonorensis.

ثم استخدام المستخلص الخالى من الخلايا المنتج من العزلات البكتيرية فى تجربة تغذية لفئران التجارب لمعرفة مدى أمان وسلامة هذه المواد فى استخدامها بالنسبة للإنسان. **الكلمات الدالة** : المضادات الحيوية – العزل والتعريف - تغذية حيوانات التجارب – الباسيلس.





igure (1): Neighbor joining tree showing the estimated phylogenetic relationships of A6 and A13 and nearest members of *Bacillus* strains. Bootstrap values are shown as • percentages of 1000 replicates. Bar: 0.1% sequence divergence.