

In Vitro Evaluation of Antibacterial Potential of *Cerastes Viper* Venom against Gram-Positive and Gram-Negative Bacterial Strains

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ABSTRACT

Background: Venoms of the family *Viperidae* in particular are complex mixture of many substances. They include many different proteins that possess significant bactericidal inhibition.

Objectives: The aim of the current work was to investigate the antibacterial profile of Sahara sand viper (*Cerastes vipera*) snake venom against both gram negative *Pseudomonas aeruginosa* and gram-positive bacteria *Bacillus cereus*.

Materials and Methods: 1. Identification of the venom was performed using assessment of total protein, determination of the venom protein by SDS-PAGE electrophoresis and determination of the half lethal dose (LD50) of venom. 2. Antibacterial activities of *Cerastes vipera* venom was examined by using disc diffusion method. Microdilution method was used to determine the venom's minimum inhibitory concentration (MIC) and the minimum bactericidal concentration (MBC). 3. The interaction between venom and bacteria was examined by Electron microscope.

Results: The total protein of the venom was 1.047±0.1 mg/ml. The SDS electrophoresis showed there were four clear bands of protein nearly at 25, 35, 48, 63 Kds and another band between 25, and 35 Kds. The LD50 was 18.3 µg/mouse (0.915 mg/kg body weight). Concerning the antibacterial effect of tested venom, disc diffusion test showed a ring of inhibition zone of about 12 mm in the case of *Pseudomonas aeruginosa* and of about 0.8 mm in the case of *Bacillus cereus*. The Microdilution test showed that the MIC of the venom for *Pseudomonas aeruginosa* was 12.5 mg/ml and that for *Bacillus cereus* was 25 mg/ml. The electron microscopic examination supported the two previous tests, as it observed the presence of elongated, malformed, melted, no-membranous and lysed bacterial cells when treated by the venom.

Conclusion: It could be concluded that *Cerastes vipera* venom is a potentially weak bactericidal, but can inhibit antibiotic-resistant bacteria as *Pseudomonas aeruginosa*.

Keywords: *Cerastes vipera* venom, bactericidal, *Bacillus cereus*, *Pseudomonas aeruginosa*, MIC, MBC, Electron microscopic examination.

INTRODUCTION

Egypt as a desert land comprises different species of venomous animals including snakes. In Egypt, 1000–10,000 incidences of snake's envenomation per year with about 11–100 death cases annually ⁽¹⁾. *Cerastes vipera* is one of the most venomous snakes of Egypt. Sahara Sand Viper (*Cerastes vipera*) is a venomous viper species endemic to the deserts of North Africa and the Sinai Peninsula ⁽²⁾.

Antimicrobial resistance is an increasingly serious worldwide threat associated with millions of illnesses and deaths annually over the world. Antimicrobial resistance is predicted to cause ten million deaths in 2050 ⁽³⁾. Given its vast ability as the source of therapeutics, it was found that viper venom has antimicrobial activity against gram-positive and gram-negative bacteria ⁽⁴⁾. The venom antimicrobial activity was due to enzymes such as L-amino acid oxidase ⁽⁵⁾.

The antimicrobial potential of snake venom enzymes is widely investigated. PLA2 and L-amino acid oxidases (LAAO1 and LAAO2) are among the best studied enzymes associated with antibacterial properties ⁽⁶⁾. Both PLA2 and LAAO are usually

responsible for the major toxic effects during snake envenomation.

The aim of the current work was to investigate the antibacterial profile of Sahara sand viper (*Cerastes vipera*) snake venom against both gram negative *Pseudomonas aeruginosa* and gram-positive bacteria *Bacillus cereus*.

MATERIALS AND METHODS

Ethical approval:

The study was approved by the Ethics Board of Al-Azhar University.

Snake venom collection: Lyophilized pure venom of Sahara sand viper (*Cerastes vipera*) was obtained from ANDI COE, VACSERA, where venom was stored in brown vial at refrigerator (4°C) till used.

Strains: *Pseudomonas aeruginosa* was obtained from American Type Culture Collection number 27853(ATCC). *Bacillus cereus* was obtained from the microbial chemistry department, National Research Center.

Inoculums Preparation: A loopful from the overnight growth was transferred to 50 ml Tryptone Soya Broth (TSB) medium and was incubated at 37°C with vigorous shaking (150 rpm) for 18 h; cell growth was checked purity by Gram stain.

Total Protein: Total protein content was determined at center of excellence in anti-venom research EGYVAC, VACSERA according to **Lowry et al.** (7).

Determination of the venom protein by SDS-PAGE electrophoresis. The venom protein activity was also assayed by electrophoresis in SDS-PAGE **Laemmli** (8) (12% acrylamide gel under reducing condition) with co-polymerized gelatin. The venom samples (10 mg) diluted in 0.05 M TRIS-glycine buffer, pH 6.8, containing 10% glycerol, 0.025 M SDS, and 0.5 mg/ml phenol red were directly applied on the gel. Soon after the run at 150 V for about 3 hours at 4°C, the gel was then (a) incubated at 37°C in 0.1 M glycine-NaOH, pH 9.5, for 5 hours, and (b) developed for proteolytic activity with 0.1% Amido Black in methanol: acetic acid: water (3:1:6; v:v:v). The gels were de-stained with methanol: acetic acid: water (3:1:6; v:v:v) until the clear zones of protein could be seen.

Determination of LD50 of venom:

The LD50 of the venom was determined according to the method of British Pharmacopeia (2000), **Reed and Muench** (9), using male albino Swiss mice 20 grams. Ascending concentration of 5 dose levels of the freshly prepared venom solution in normal saline with an increasing factor 1.2, starting by a dose which kills approximately 0-20 % of the animals and ending by a dose which kills approximately 80-100 % of the injected animals. Each dose level was tested in 4 mice, and all injections were given intravenously.

Disc diffusion method: The surface of the plates was uniformly covered with the bacterial growth for *Pseudomonas aeruginosa* and *Bacillus cereus* which re-suspended and adjusted turbidity to 0.5 **McFarland** (10). The sterile filter paper discs were impregnated in venom with concentration of 100mg venom /1ml saline.

- Disc of antibiotics like erythromycin (15 µg Oxoid lot no. 1876568), Ampicillin (10 µg Oxoid lot no. 1843425), Amikacin (30 µg Oxoid lot no.2288626) and Ciprofloxacin (5 µg Oxoid lot no.2280856) as positive control. The discs were placed on the surface of the plates suitably spaced apart, and then the plate was incubated at 37°C for 18 hrs. The plate was examined for the presence of zone of inhibition for bacterial growth around the disc. The zone of inhibition was indicative the degree of organism sensitivity (11).

The minimum inhibitory concentration (MIC) and

minimum bactericidal concentration (MBC):

The bacterial growth was re-suspended and adjusted turbidity to 0.5 **McFarland** (10) using spectrophotometer (Perkin Elmer-EZ 301-German) for *Pseudomonas aeruginosa* and *Bacillus cereus* according to **Andrews** (12), 100 mg venom /1ml saline. 100µl of standard **McFarland**(10) bacterial growth of each strain was pipette into each well of a microtiter plate (96 Well CELLSTAR®, Greiner Bio-One GmbH). Wells no. 1&2 were positive control and Wells no. 11&12 were negative control. 100µl of antibiotic was pipette into wells no. 3&4 as positive control. The plate 1 for *Bacillus cereus*, a serial dilution (1/2 – 1/64) from concentration of 100mg venom /1ml saline was added from well no. 3 to well number 10. The plate 2 for *Pseudomonas aeruginosa*, a serial dilution (1/2 – 1/64) from concentration of 100mg venom /1ml saline was added from well no. 3 to well number 10. The plates were incubated at 37°C for overnight. The plates were read on ELISA reader (Biotek- ELX800- USA) at wavelength 570 nm.

Electron microscopic examination: Preparation of the sample: 1-Bacteria:

A loopful from the overnight growth was transferred to 50 ml Tryptone Soya Broth (TSB) medium and was incubated at 37°C with vigorous shaking (150 rpm) for 18 h; cell growth was checked purity by Gram stain. **2-Specimen:** The Prepared bacteria was mixed with the MIC of the venom and incubated at 37°C for 48 hours.

Negative Stain Biological Specimen Preparation (Non sonicator):

- 1) Use a 10 ml disposable pipet to resuspend the cells by pipetting up and down.
- 2) Place 2-5 ul drops of specimen onto sheet of Parafilm.
- 3) Make EM grids (carbon-coated 400-mesh copper grids) directly off on specimen.
- 4) Use filter paper to wick away specimen drop.
- 5) Place grid on drop of filtered 2% PTA stain, pH 7.0, and let stain for 30-60 sec.
- 6) Use filter paper to wick away stain and place grids, specimen-side up in specimen petri dish.

RESULTS

Experiment I: venom identification

Assessment of Total protein:

Table 1 showed the value of total protein of *Cerastes vipera* venom which equal 1.047±0.1 mg/ml.

Table 1: Total protein of *Cerastes vipera* venom

Protein concentration	Warburg and Christian, 1942 (12)	Up to 2 mg/ml	1.047±0.1 mg/ml
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Biochemical characterization by SDS page electrophoresis

protein nearly at 25, 35, 48, 63, Kds and another band bet ween 25 and 35Kds.

From fig. 1 we can represent four clear bands of

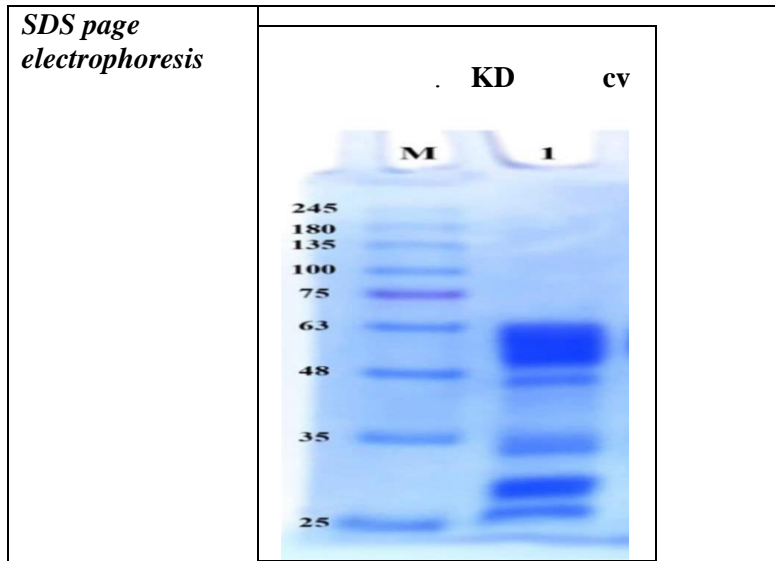


Fig.1. SDS page electrophoresis for *Cerastes vipera* venom

The assessment of half lethal dose (LD50):

Table 2 and figure 2 showed the approximate half lethal dose (LD50) of *Cerastes vipera* venom that assessed according to Reed and Muench, (1938) ⁽¹¹⁾.

LD50 = 18.3 µg/mouse (0.915 mg/kg body weight).

Table 2: Lethality of *Cerastes vipera* venom injected I/V in Swiss-albino mice and estimated value of LD₅₀

Dose (µg/mouse)	Accumulated Values					Mortality Ratio (%)
	Died	Survived	Died	Survived		
11.5	0	5	0	15		00.0%
13.8	1	4	1	10		0.9%
16.6	2	3	3	6		33.3%
20	3	2	6	3		66.6%
24	4	1	10	1		90.9%
28.8	5	0	15	0		100%

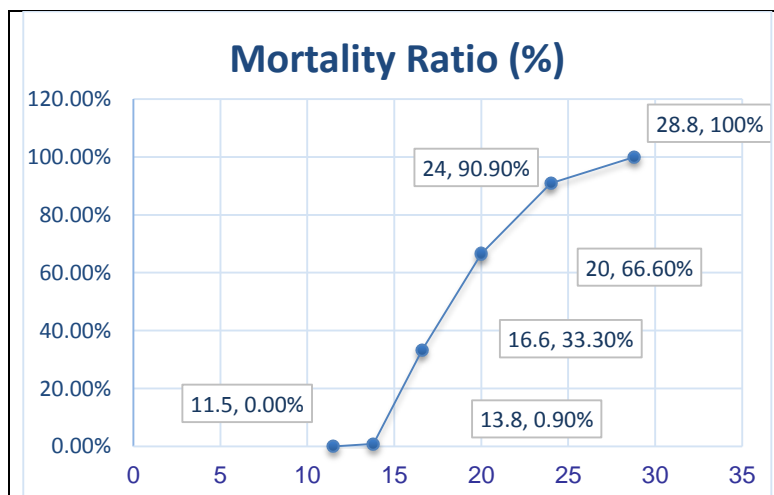


Fig. 2: Mortality curve of Swiss Albino mice administrated different doses of *Cerastes vipera* venom

Experiment II: Antibacterial effect of *Cerastes vipera* venom:

A) Model of gram-negative bacteria: *Pseudomonas aeruginosa*

1. Disc diffusion

The disc diffusion method was used to assess the antibacterial effect of *Cerastes vipera* venom compared with the same standard antibiotics; Erythromycin (E), Ampicillin (AMP), Ciprofloxacin (CIP) and Amikacin (AK), against *Pseudomonas aeruginosa* (Table 3).

Table 3: In vitro antibacterial activity of *Cerastes vipera* crude venom (20 mg/ml) against *pseudomonas aeruginosa* strain using disc diffusion method and compared to four antibiotics (stander disc). (Mean ± S.D) of inhibition zone in mm.

Antibiotics / Venom	Inhibition zone
<i>Cerastes vipera</i> venom (20 mg/ml)	(12± 0.1) mm
Ampicillin (10 µg/ml)	(0.0±0.0) mm
Erythromycin (15 µg/ml)	(0.0±0.0) mm
Ciprofloxacin (5 µg/ml)	(30±0.11) mm
Amikacin (30 µg/ml)	(20±0.09) mm

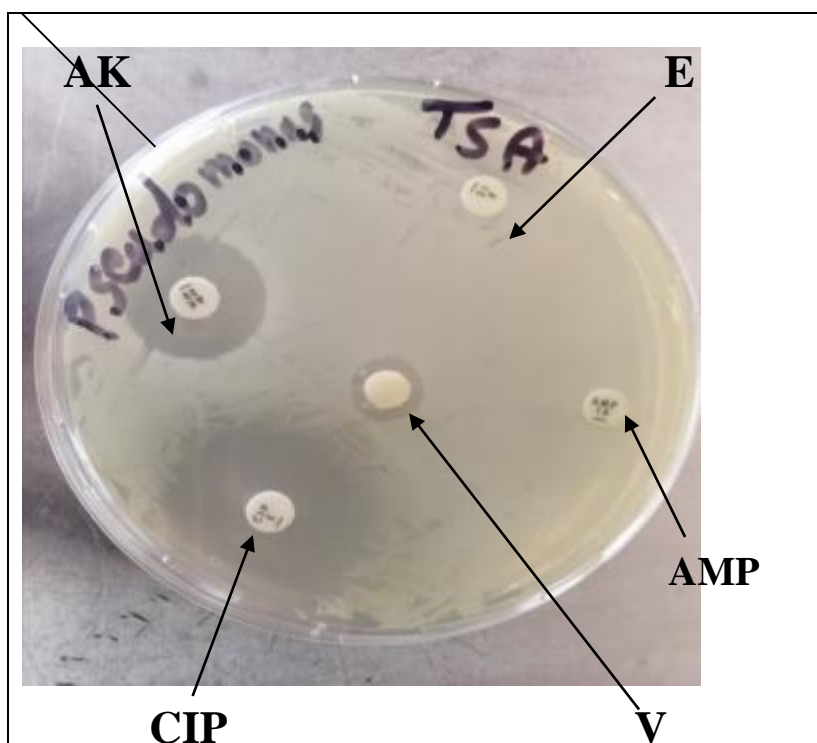


Fig.3: Antibacterial effect of *Cerastes vipera* venom (V) against *pseudomonas aeruginosa* in compared with four antibiotics; Ampicillin (AMP), Erythromycin (E), Ciprofloxacin (CIP) and Amikacin (AK)

There was a highly effective antibacterial effect of Ciprofloxacin antibiotic with the *pseudomonas aeruginosa* bacterial growth producing a wide ring of inhibition zone average; 30 mm, Amikacin antibiotic produced intermediate ring of inhibition zone about 20mm, However the *Cerastes vipera* venom produced low ring of inhibition zone of about 12mm and Ampicillin and Erythromycin produced no inhibition zone (resistance) as shown in (Table 3&Fig. 3).

Determination of Minimum Inhibitory Concentration (MIC)

From table 4, and figure 4, It can be observed that the minimum Inhibitory concentration (MIC) of *Cerastes vipera* venom in the case of *Pseudomonas aeruginosa* is 12.5mg/ml which gave inhibition value equal 0.68±0.011 and it is higher than the MIC of tylosin antibiotic for the same bacteria that is 10mg/ml which gave

inhibition value equal 2.52 ± 0.16 .

Table 4: Determination of Minimum Inhibitory Concentration (MIC) for *Pseudomonas aeruginosa*

<i>Pseudomonas aeruginosa</i>	Tylosin	Venom	control +	control-
			0.5 Mcf	media
MIC value (concentration)	10 mg/ml	12.5 mg/ml	bacteria	
Inhibition value	2.52 ± 0.16	0.68 ± 0.011	1.72 ± 0.045	0.24 ± 0.04 0.37 ± 0.035
MIC values are given as mean of five replicates (n = 8), the bacterial inoculums per plate contained 3.2×10^8 cfu/mL forming units/well, Control (0.5 MQ) (bacterial inoculums); venom breaking points (MICs 12.5 mg/mL) against <i>P. aeruginosa</i> bacteria, Results is expressed by mean \pm SD				

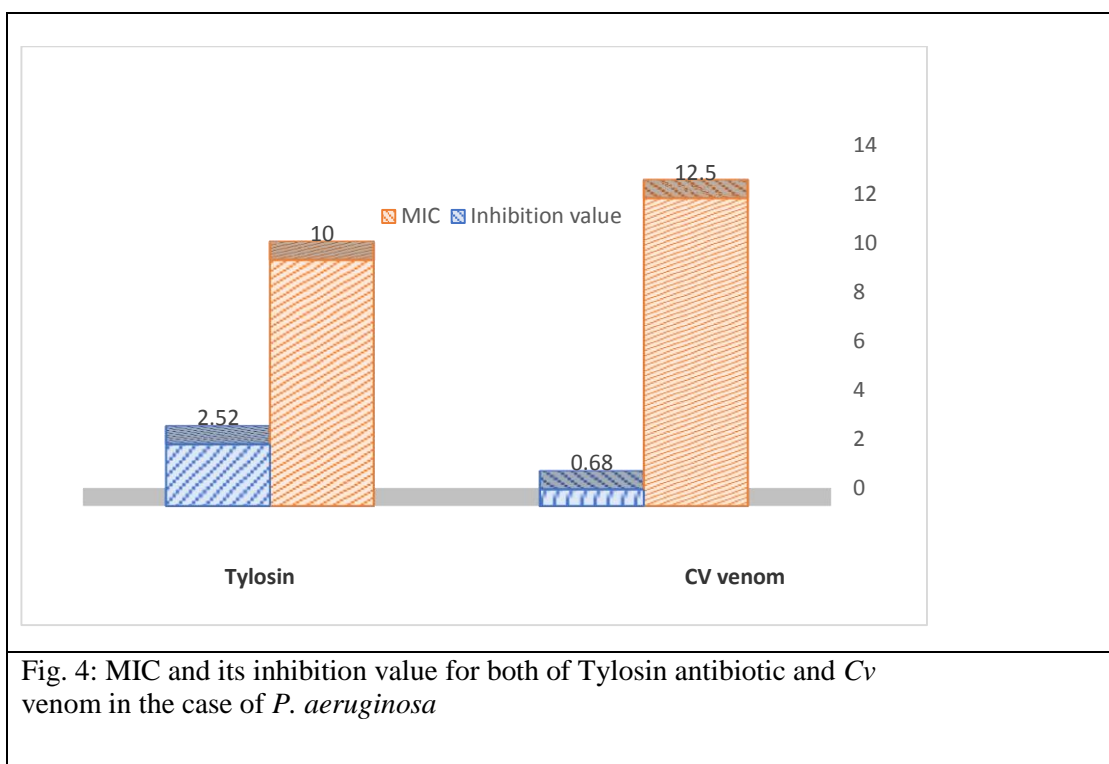


Fig. 4: MIC and its inhibition value for both of Tylosin antibiotic and Cv venom in the case of *P. aeruginosa*

2. Determination of Minimum Bactericidal Concentration (MBC)

From table 5, and figure 5, It can be observed that the minimum bactericidal concentration (MBC) of *Cerastes vipera* venom in the case of *Pseudomonas aeruginosa* is 25mg/ml which gave inhibition value equal 0.82 ± 0.009 and it is higher than the MBC of tylosin antibiotic for the same bacteria that is 20 mg/ml which gave inhibition value equal 3.5 ± 0.18

Table 5: Determination of the Minimum Bactericidal Concentration (MBC) of Cv for (*Pseudomonas aeruginosa*)

<i>Pseudomonas aeruginosa</i>	tylosin	cv	control +	control-
			0.5 Mcf bacter	media
MBC value (concentration)	20 mg/ml	25mg/ml		
Inhibition value	3.5 ± 0.18	0.82 ± 0.009	1.72 ± 0.045	0.37 ± 0.035

values are given as mean of five replicates (n = 8), the bacterial inoculums per plate contained 3.2×10^8 cfu/mL forming units/well, Control (0.5 MQ). *The toxin breaking points (MBCs 25.0 mg/mL) against *Pseudomonas aeruginosa* bacteria after 24 h incubation

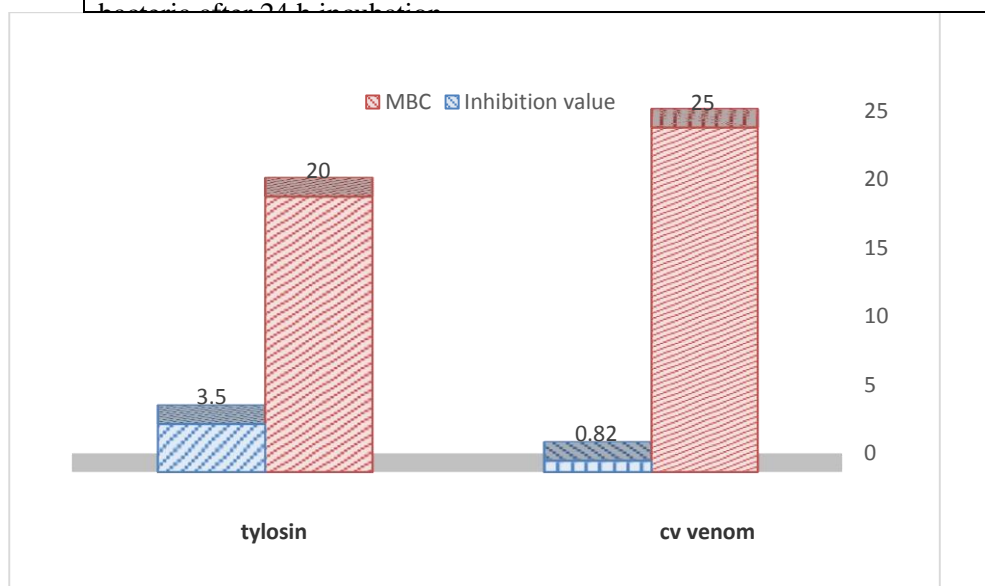


Fig. 5: MBC and its inhibition value for both of Tylosin antibiotic and Cv venom in the case of *Pseudomonas aeruginosa*

B) Model of gram-positive bacteria: *Bacillus cereus*

1. Disc diffusion

The disc diffusion method was used to assess the antibacterial effect of *Cerastes vipera* venom compared with different standard antibiotics; Erythromycin (E), Ampicillin (AMP), Ciprofloxacin (CIP) and Amikacin (AK), against *Bacillus cereus* Table 6 and figure 6.

Table 6: *In vitro* antibacterial activity of *Cerastes vipera* crude venom (20 mg/ml) against *bacillus cereus* strain using disc diffusion method and compared to four antibiotics (stander disc). (Mean ± S.D) of inhibition zone in mm.

Antibiotics / Venom	Inhibition zone/mm
<i>Cerastes vipera</i> venom (20 mg/ml)	8± 0.2
Ampicillin (10 µg/ml)	20±0.12
Erythromycin (15 µg/ml)	32±0.09
Ciprofloxacin (5 µg/ml)	40±0.11
Amikacin (30 µg/ml)	28±0.09

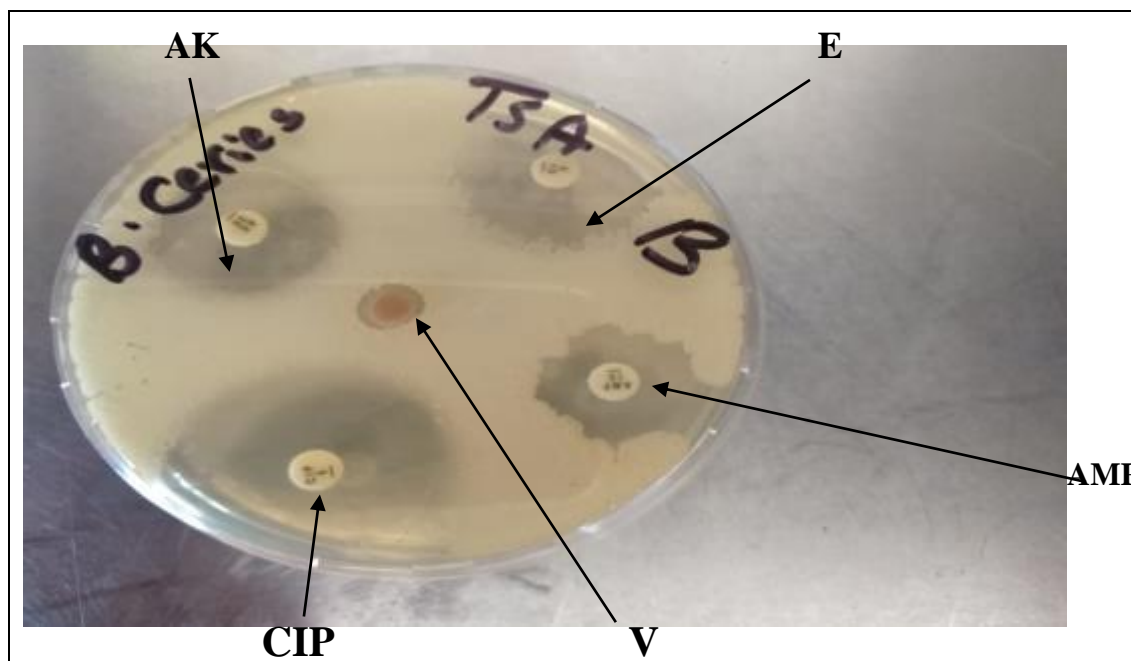


Fig.6: Antibacterial effect of *Cerastes vipera* venom (V) against *Bacillus cereus* in compared with four antibiotics; Ampicillin (AMP), Erythromycin (E), Ciprofloxacin (CIP) and Amikacin (AK)

There was a highly effective antibacterial effect of Ciprofloxacin, Erythromycin and Amikacin antibiotics produced a wide ring of inhibition zone about 40, 32, 28 mm respectively, while the Ampicillin antibiotic produced intermediate inhibition zone average; 20 mm and the *Cerastes vipera* venom produced low ring of inhibition zone of about 8 mm, all with the *Bacillus cereus* bacterial growth (Table6 & Fig.6).

2. Determination of the Minimum Inhibitory Concentration (MIC)

From table 7, and figure 7, It can be observed that the minimum inhibitory concentration (MIC) of *Cerastes vipera* venom in the case of *bacillus cereus* is 25mg/ml which give inhibition value equal 0.8 ± 0.062 and it is higher than the MIC of oxytetracycline antibiotic for the same bacteria that is 5mg/ml which give inhibition value equal 0.66 ± 0.028 .

Table 7: Determination of Minimum Inhibitory Concentration (MIC) for *Bacillus cereus*

<i>Bacillus cereus</i>	oxytetracycline	venom	control	Control-
MIC value (concentration)	5 mg/ml	25 mg/ml	0.5 Mcf bacteria	media
Inhibition value	0.66 ± 0.028	0.8 ± 0.062	1.83 ± 0.032	0.42 ± 0.045
MIC values are given as mean of five replicates (n = 8), the bacterial inoculums per plate contained 3.2×10^8 cfu/mL forming units/well, Control (0.5 MQ) (bacterial inoculums); venom breaking points (MICs 25 mg/mL) against <i>P. aeruginosa</i> bacteria. Results represented by mean \pm SD				

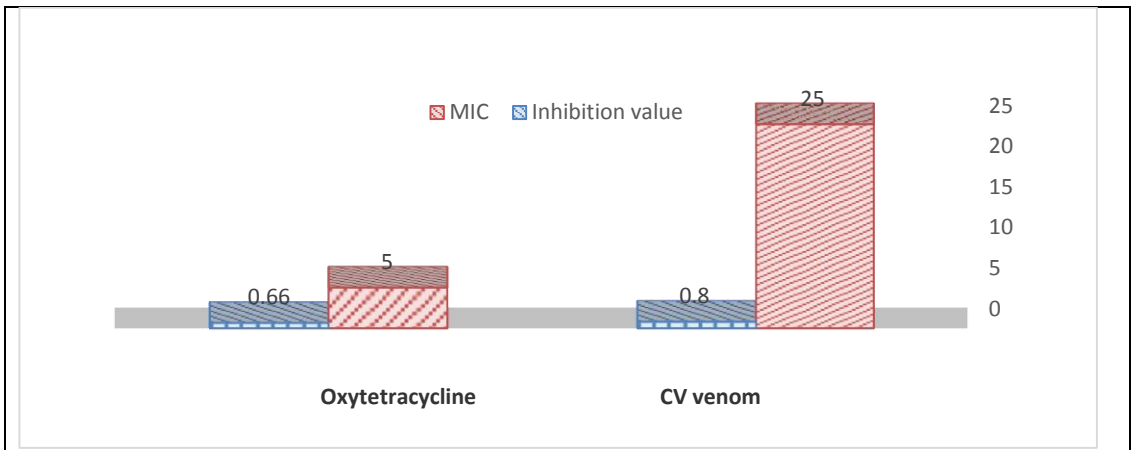


Fig. 7: MIC and its inhibition value for both of oxytetracycline antibiotic and Cv venom in the case of *B. cereus*

3. Determination of the Minimum Bactericidal Concentration (MBC)

From table 8, and figure 8, It can be observed that the minimum Bactericidal concentration (MBC) of *Cerastes vipera* venom in the case of *Bacillus cereus* is 50mg/ml which give inhibition value equal 0.87 ± 0.07 and it is higher than the MBC of oxytetracycline antibiotic for the same bacteria that is 10mg/ml which give inhibition value equal 0.80 ± 0.03 .

Table 8: Determination of the Minimum Bactericidal Concentration (MBC) for *Bacillus cereus*

<i>Bacillus cereus</i>	oxytetracycline	venom	control	control-
MBC value (concentration)	10 mg/ml	50mg/ml	0.5 Mcf bacteria	media
Inhibition value	0.80 ± 0.03	0.87 ± 0.07	1.83 ± 0.032	0.42 ± 0.045

values are given as mean of five replicates (n = 8), the bacterial inoculums per plate contained 3.2×10^8 cfu/mL forming units/well, Control (0.5 MQ) (bacterial inoculums); The toxin breaking points (MBCs 50 mg/mL) against *Bacillus cereus* bacteria after 24 h incubation.

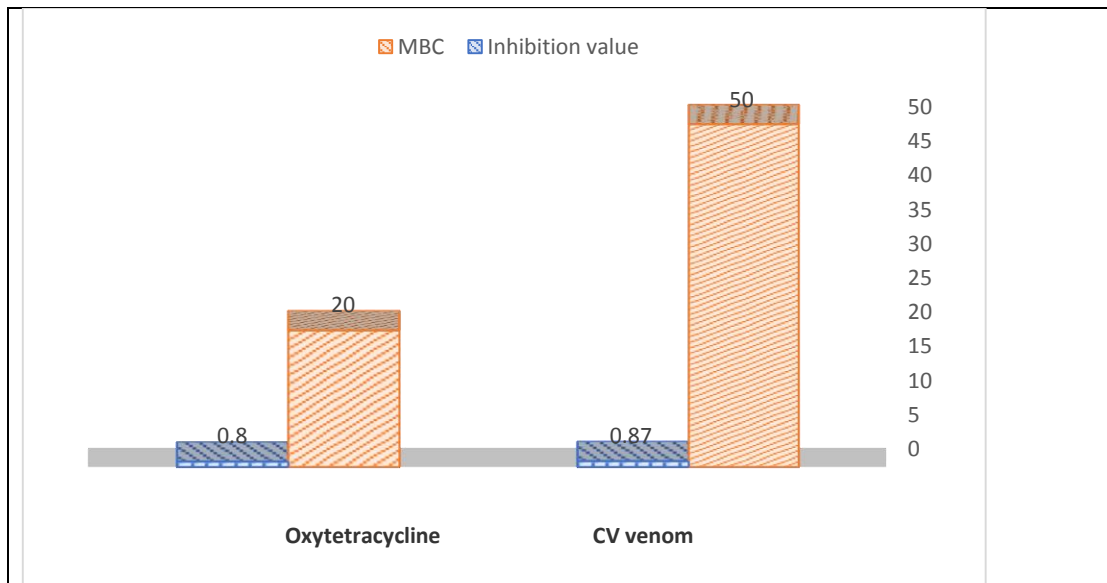


Fig. 8: MBC and its inhibition value for both of oxytetracycline antibiotic and Cv venom in the case of *Bacillus cereus*

Experiment III: Electron microscope examination

A) Model of gram-negative bacteria: *Pseudomonas aeruginosa*

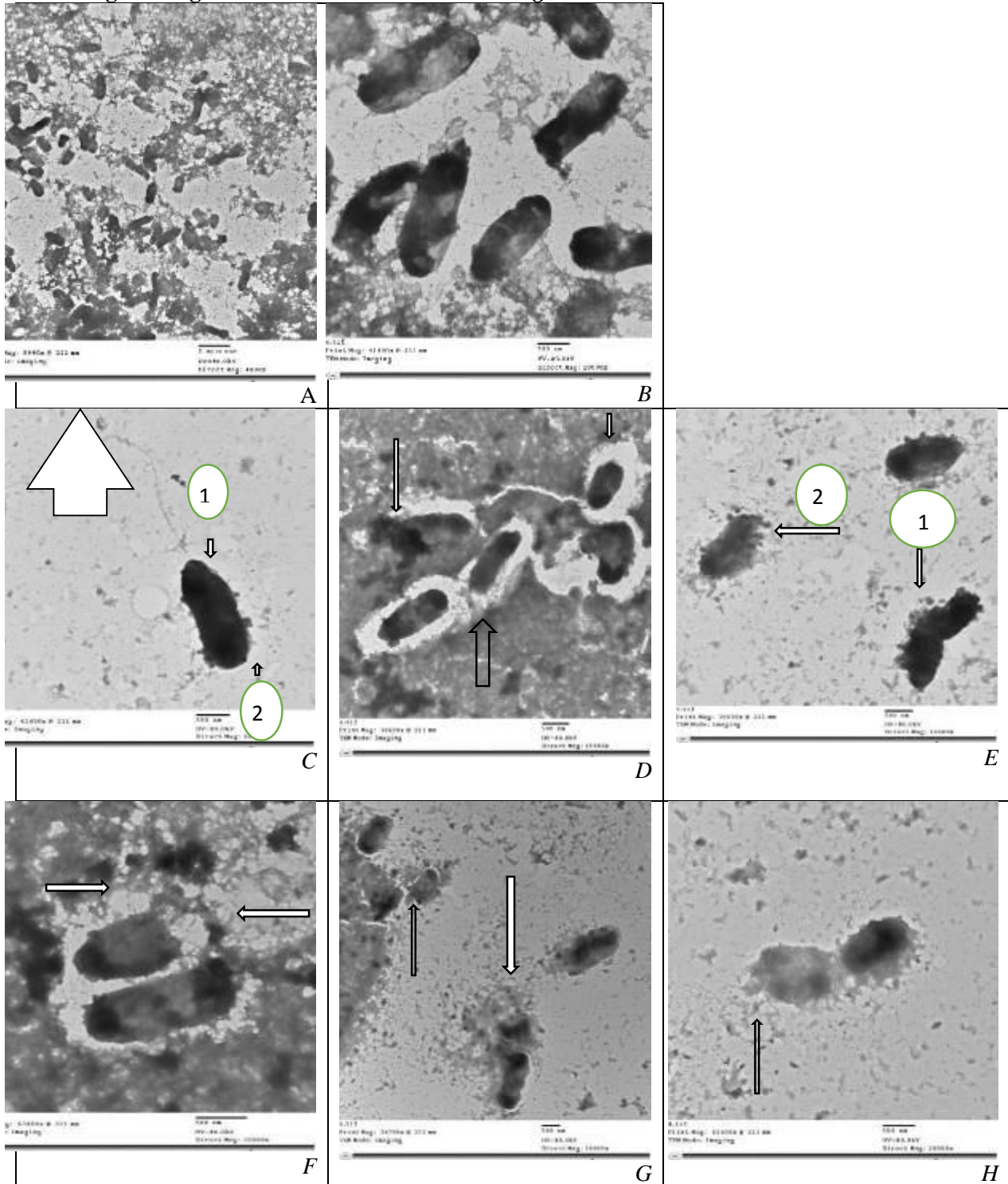


Fig. 9: TEM of untreated (control) and treated *P. aeruginosa* with *C. vipera* venom

The untreated (control) cells of *P. aeruginosa* appeared bacilli and without visible abnormalities. The formation of a septum was also observed, which indicated that active proliferation was occurring. The cells were separated from each other and displayed a smooth and intact surface with undamaged cell walls (Figure 9A, 9B). After incubation with the MIC of the *C. vipera* venom for 48 h, the bacteria appeared elongated (fig. 9C1, 9E1). In addition, the cells appeared aggregated compared with the control, some of these melted (fig. 9D). The treated bacterial cells revealed variation in the cell wall thickness and alteration of the shape, which appeared elongated and malformed compared with the control (fig. 9D). Additionally, large spherical non-membrane-enclosed bodies with a similar electron density to that of the septal murein layer were observed in it (Fig. 9F, 9G). There were lysed cells with severely damaged and depleted cell content and with a lamellar form of the cytoplasm were also notice (fig. 9H).

B) Model of gram-positive bacteria: *Bacillus cereus*

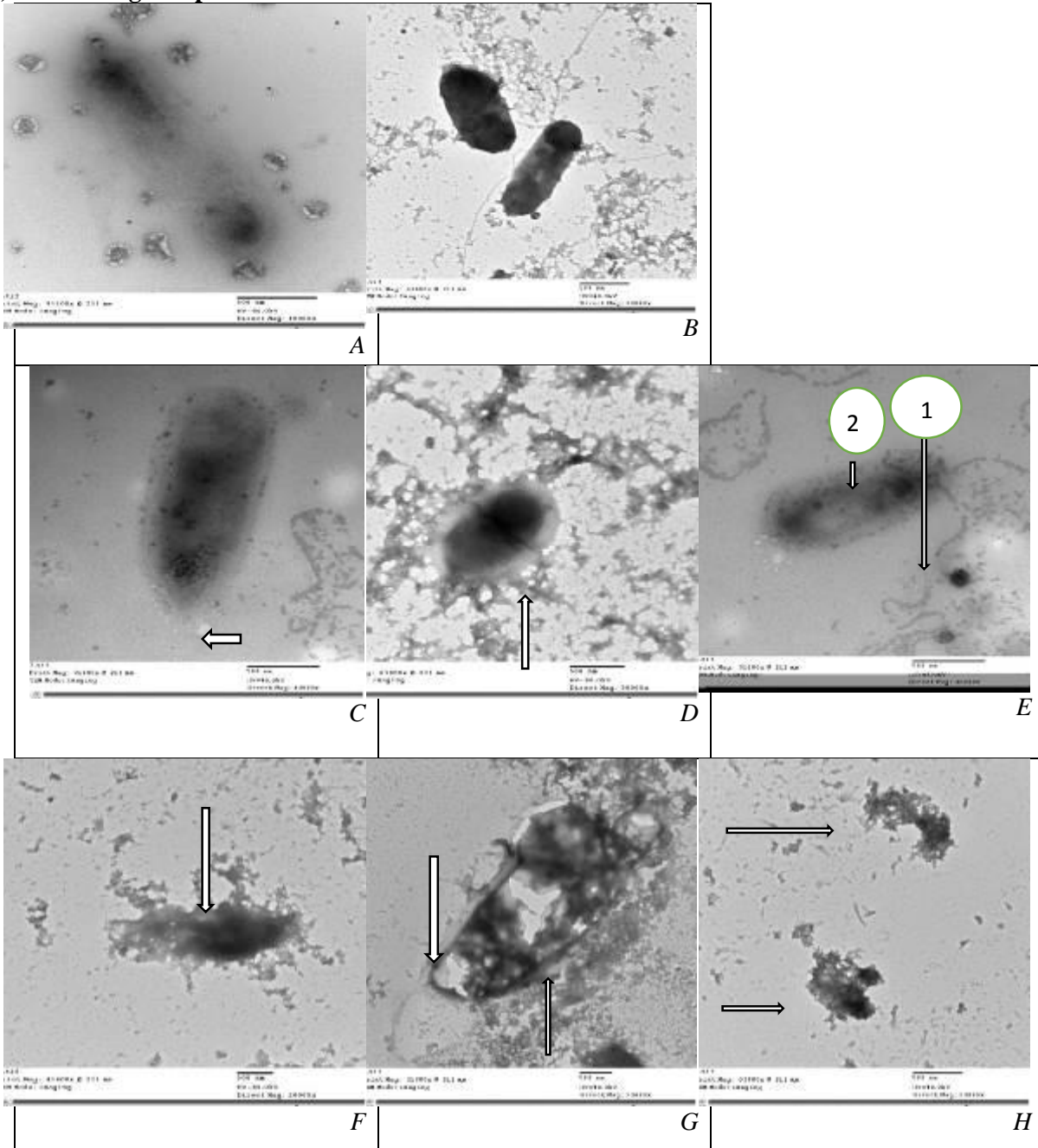


Fig. 10: TEM of untreated (control) and treated *B. cereus* with *C. vipera* venom

The untreated (control) cells of *B. cereus* appeared bacilli and without visible abnormalities. The formation of a septum was also observed, which indicated that active proliferation was occurring. The cells were separated from each other and displayed a smooth and intact surface with undamaged cell walls (Figure 10A, 10B). After incubation with the MIC of the *C. vipera* venom for 48 h, the bacteria appeared elongated (fig. 10C). In addition, the treated bacterial cells revealed variation in the cell wall thickness and it became more transparent compared with the control (fig 10C, 10E2). Additionally, large spherical non- membrane-enclosed bodies with a similar electron density to that of the septal murein layer were observed in it (Fig. 10E1). The periplasmic space appeared separated, suggesting detachment of the cell wall from the plasma membrane (fig. 10D), and some had holes in their cell wall (Fig. 10 G). The components of the bacterial cell envelope were deformed and scattered from their original form (Fig. 10E1, 10H). At certain locations of the cells, mostly the septal and polar regions, a draining out of the intracellular contents due to rupture of the cell envelope was noted. Null cells were found (Fig. 10 H) and many cells without membranes (Fig. 10 F).

DISCUSSIONS

Antibiotic-resistant bacteria strains are of the top leading causes of death worldwide (4). According to the WHO, they are one of the ten leading causes of death worldwide (13-14). Moreover, antimicrobial resistance is considered to be one of the greatest threats to human health globally (15). In this context, the discovery of new antimicrobial agents is crucial, and the search for new compounds has led to the exploration of various natural sources, such as plant extracts (16-18), bee venom (19), spider venom (20) or snake venom (21). Several studies proved the antibacterial effect of snake venoms (22).

In the present study, first, the identification of *Cerastes vipera* crude venom evaluated assessment of total protein, Biochemical characterization by SDS page electrophoresis and the assessment of half lethal dose (LD50). The value of total protein of *Cerastes vipera* venom was equal 1.047 ± 0.1 mg/ml. This result is in agreement with **Abd El – Aal et al.** (23) who reported that the Total protein of *Cerastes vipera* venom is 0.927 mg/ml. In order to make a Comparison between the total protein of *Cerastes vipera* venom and some other species of *Viperidae* family, it was mentioned that 1mg of lyophilized venom of *Macrovipera mauritanica* contains 1.0 mg of proteins and about 0.987 mg of proteins for *Cerastes cerastes* venom (24).

Examination of CV venom using SDS-PAGE electrophoresis revealed that there are four clearbands of protein nearly at 25, 35, 48, 63 Kds and another band between 25, and 35 Kds Our result nearly agrees with that of **Labib et al.** (25) who observed that there are six major bands. The assessment of LD50 of the *Cerastes vipera* crude venom was recorded 18.3 µg/20 mouse (0.915 mg/kg body weight). This result agrees with that obtained by **Mohamed et al.** (26) and **Hassan et al.** (27) who reported that the *C. vipera* venom is lethal in animals (rabbits LD50 4 mg/kg; mice LD50 1 mg/kg; rats LD5 1.3 mg/kg, and doesn't match with that of **Abdel-Aty et al.** (28) who reported that the LD50 of Cv crude venom was recorded at 10 µg/ 21 g mouse.

Second, the antibacterial activity of *Cerastes vipera* crude venom against the selective bacterial strains was evaluated by disc diffusion technique, the venom had weak effect against Gram-positive bacteria *Bacillus cereus* (with inhibition zone 8 ± 0.2 mm) in compared with standard antibiotics Ampicillin (20 ± 0.12 mm), Erythromycin (32 ± 0.09 mm), Ciprofloxacin (40 ± 0.11 mm) and Amikacin (28 ± 0.09 mm) that were highly effective antibacterial. Also, the venom showed acceptable effectiveness against Gram-negative bacteria *Pseudomonas aeruginosa* (with inhibition zone 12 ± 0.1 mm) in compared with the same standard antibiotics, as this type of bacteria was resistant to Ampicillin and Erythromycin (both with no inhibition

zone). Also, *Pseudomonas aeruginosa* classified as a member of ESKAPE group that reference to their ability to escape the effects of commonly used antibiotics through evolutionarily developed mechanisms (29).

The result was nearly agreed with **Salama et al.** (30) who reported that L-amino acid oxidase isolated from *Cerastes vipera* snake venom displayed bactericidal effect and showed inhibition zone 15 ± 0.2 mm against *pseudomonas aeruginosa* and showed low inhibition zone 8 ± 0.2 mm against *Bacillus cereus*. In contrast, **Shebl et al.** (31) reported that the venom of both *Cerastes cerastes* and *Vipera lebetina* showed no inhibition zone against *pseudomonas aeruginosa*, this difference due to the variation in *Viperidae* family species. On the other hand, antibacterial effect was evaluated by assessment the minimum inhibitory concentration (MIC) and minimum bactericidal concentration (MBC).

The result showed The MIC in the case of *Bacillus cereus* was 25mg/ml and MBC was 50mg/ml, whereas the MIC in the case of *Pseudomonas aeruginosa* was 12.5mg/ml and MBC was 25mg/ml. Our results showed very high level of both MIC and MBC in compared with those of **Walaa H. Salama et al.** (30) who reported that the minimum inhibitory concentrations (MIC) of the *Cerastes vipera* L-amino acid oxidase enzyme against the gram-positive strain *S. aureus* was found to be 20 µg/ml, while MIC values of this enzyme against *P. aeruginosa* was found to be 40 µg/ml, with consideration that L-amino acid oxidase represent 9% of *cerastes vipera* venom, according to **Amine Bazaa et al.** (32) this difference due to difference between crud venom and effective pure fraction.

Third, the electron microscopic examination for the untreated (control) cells of both *P. aeruginosa* and *B. cereus* appeared bacilli and without visible abnormalities. The surface of the bacterial cells appeared smooth with undamaged cell walls. These results were similar to that of **Ramar Perumal Samy et al.** (33) who noticed that the untreated bacteria appeared with smooth membrane structure when examined by electron microscope, and agree with **Zhang et al.** (34) who concluded that the surface of bacteria was smooth with a distinct boundary between the cells. When tested *P. aeruginosa* treated by the Cv venom and examined by TEM, it appeared in multiple shapes such as variation in the cell wall thickness, aggregated, melted, elongated, malformed, lysed cells with severely damaged and depleted cell content.

On the other hand, *Bacillus cereus* treated by the venom also appeared more Transparent, some became polar, the periplasmic space appeared separated (suggesting detachment of the cell wall from the plasma membrane), large spherical non-membrane-enclosed bodies appeared, holes in the cell wall,

draining out of the intracellular contents and null cells were found. These results were similar to that of **Samy et al.** (33) who reported that bacteria appeared to be losing cell contents and losing of the membrane structure. Also, our results match with **Zhang et al.** (34) who illustrated that the membrane of bacteria treated with antibacterial protein was largely damaged, resulting in the release of the cell content upon cell wall disruption, with noticeable gaps between the cell membrane and the cytoplasm. **Luyang et al.** (35), also showed that some venom peptide derivatives had make structural changes and cracks in the cell walls and membranes of the bacteria, causing a release of cell content upon cell wall disruption.

CONCLUSION

It could be concluded that *Cerastes vipera* venom is a potentially weak bactericidal, but can inhibit antibiotic-resistant bacteria as *Pseudomonas aeruginosa*.

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