CHARACTERIZATION OF GRAM-NEGATIVE BACTERIA ISOLATED FROM LUNGS OF CAMEL

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ABSTRACT

Out of 308 samples 205 from apparently healthy and 103 diseased lung of slaughtered camels. The total number of positive samples was 203 (65.91%) from which 124 (60.49%) from apparently healthy and 79 (76.70%) from pneumonic lung of camels.

The total number of isolates was 302 isolates from which 174 isolates from apparently healthy and 128 isolates from pneumonic lung of camels. The total number of Gram negative isolates 170 (56.29%) from which isolates were 94 (54.02%) and 76 (59.38%) in apparently healthy and diseased lungs of camel respectively as shown in table (3).

On the other hand the main predominant single Gram negative isolates were E. coli 28 samples (26.92%), Pseudomonas aeruginosa 9 samples (8.65%), Citrobacter freundii 7 samples (6.73%), Klebsiella pneumoniae 6 samples (5.77%), Citrobacter spp. 5samples (4.81%), Klebsiella oxytoca 3 samples (2.89%), Mannheimia haemolytica and Yersinia enterocolitica 2 samples (1.92%), Klebsiella ozaenae and Pasteurella multocida one sample (0.96%). Serological identification from apparently and pneumonic lung of slaughtered camels was 44 isolates of E. coli including 29 isolates from apparently healthy as E. coli O_{111} , E. coli O_{115} , E.coli O_{119} E. coli O_{146} and E. coli O_{166} and 20 isolates from pneumonic lung including E. coli O_{55} , E. coli O_{111} E. coli O_{115} , E. coli O_{119} , E. coli O_{125} , E. coli O_{146} and E. coli O_{166} .

Regarding to the pathogenicity of gram negative isolates in experimentally infected mice by 0.1 ml (5 x 10^8 cFm. ml). It was fatal to mice with varying pathogenicity, Klebsiella pneumoniae, E. coli O_{55} , E. coli O_{115} , Pseudomonas aerouginosa, Pasteurella multocida and Mannheimia haemolytica were 100%. Meanwhile the lowest pathogenic bacteria were E. coli O_{119} and Proteus vulgaris were 20%. But citrobacter spp. and Proteus penneri not causing deaths among mice.

Chloramphenicol, Ciprofloxacin, Enrofloxacin, Flumequine, Gentamicin and Norfloxacin against Citrobacter freundii, Citrobacter spp., E. coli, Enterobacter cloacae, K. oxytoca, K. ozaenae and K. pneumoniae. Ciprofloxacin, Doxycycline, Enrofloxacin and Norfloxacin against Mannheimia haemolytica, Pasteurella multocida, Proteus penneri, Proteus vulgaris, Ps. aeruginosa and Yersinia enterocolitica.

SDS-PAGE analysis of Klebsiellae and E. coli showed that they contained from 5 to 12 bands after staining with Commiassie brilliant blue method which molecular weight ranged from 7.218 to 170.97 KDa.

It was obvious that the molecular weight of E. coli O_{55} ranged from 134.55 to 7.2118 KDa. The molecular weight of E. coli 111 ranged from 134.55 to 32.208 KDa. The molecular weight of E. coli O_{115}

ranged from 134.55 to 33.311 KDa. The molecular weight of E. coli O_{119} ranged from 131.24 to 8.4471 KDa. The molecular weight of E. coli O_{125} ranged from 118 to 28.562 KDa. The molecular weight of E. coli O_{146} ranged from 151.11 to 11.741 KDa. The molecular weight of E. coli O_{166} ranged from 154.41 to 21.137.

On the other hand the molecular weight of Klebsiella oxytoca ranged from 157.72 to 15.859 KDa. The molecular weight of Klebsiella pneumoniae ranged from 167.66 to 22.15 KDa. The molecular weight of Klebsiella ozaenae ranged from 170.97 to 38.016 KDa.

Key Words: Klebsiella, E coli, pneumonic lung, antimicrobial agents.

INTRODUCTION

Camel plays vital socioeconomic roles and supports the survival of millions of people in Asia and Africa. It is being used as a source of protein, milk and hides as well as quiet and effective means of transport (*Chowdhary*, 1986).

Respiratory affections are the main causes of death among camel calves all over the world (*Chowdhary, 1986 and Khanna et al., 1992*).

Bacterial infection of lung is one of the main causes of pneumonia (*Thabet, 1994*). Many gram negative bacteria were isolated from pneumonic lungs as pseudomonas *spp.*, *E. coli*, Klebsiella *spp.*, *Mannheimia haemolytica, Pasteurella multocida*, and *Citrobacter spp. Thabet (1994)*, *Amany (2000)* and *Azzam and Zaki (2006)*.

The whole cell protein profiles of bacteria can be easily studied on SDS-PAGE. This powerful technique allows very high resolution of protein and has permitted the identification of multiple major proteins components of different molecular weights.

The present work was aimed to study the Gram-negative bacterial species of respiratory system of camels. For achieving the following points:

- Isolation and identification of gram negative bacteria recovered from apparently healthy and pneumonic lungs of camels.
- Serogrouping of isolated *E. coli*.
- Determination of the pathogenicity of isolates to mice.
- Detecting the antibiogram against the isolated bacteria.
- Studying the SDS-PAGE of whole cell proteins of some Klebsiellae and *E. coli* strains.

MATERIAL AND METHODS

1. MATERIALS:

1.1. Samples for isolation:

- 308 samples from imported slaughtered camels were collected from El-Basatein abattoir, Cairo-Egypt, The collected samples were 250 samples from apparently healthy and 102 samples had pathological lesions.
- All samples were kept in clean plastic ice bags in cooling container and transferred as soon as possible to the laboratory of Animal Health Research institute, Dokki, Egypt, for bacteriological examination.
- 1.2. Media used for isolation and cultivation of bacteria:
- 1.2.1. Fluid media:
- 1. 2.1.1. Nutrient broth: (Oxoid, Code CM1)

- 1. 2.2. solid media:
- 1. 2.2.1. Nutrient agar medium: (Oxoid, Code, CM3)
- 1.2.2.2. Blood agar medium: (Oxoid, Code, CM55)
- 1.2.2.3. MacConkey lactose bile salt agar: (Oxoid, Code, CM115)
- 1.2.2.4. DAS medium: (Oxoid)
- 1.2.2.7. Eosin Methylene Blue agar (EMB agar): Quinn et al. (1994)
- 1.2.3. Semisolid media: (soft agar): (Cruickshank et al. 1975)

This medium was used for the preservation of all isolates.

1.3. Media used for biochemical identification of the isolates:

All media used were prepared according to *Cruickshank et al.* (1975) and Koneman et al. (1988).

- 1.3.1. Peptone water 1%: (Oxoid, Code L37)
- 1.3.2. Glucose phosophate broth:(Oxoid, Code CM43)
- 1.3.3. Simmon's citrate agar: (Oxoid, CM115)
- 1.3.4. Urea agar base: (Oxoid, CM53)
- 1.3.5. Nutrient gelatin: (Oxoid, CM35a)
- 1.3.6. Nitrate broth: (Difco, 268)
- 1.3.7. Triple sugar iron agar medium (T.S.I): (Oxoid, Code M277)
- 1.3.8. Sugar fermentation media: (Kreig and Holt, 1984)
- 1.3.9. Phenyl alanine agar: (Cowan, 1979).
- 1.3.10. Amino acid decarboxylase test.

1.4. Stains used:

- 1.4.1.Gram's stain: It was prepared according to *Mackie and MacCarteny (1989)*.
- 1.4.2.Leishman stain: (Cruickshank et al., 1975)
- 1.5. Chemical and Reagents: It was prepared according to *Koneman et al. (1988)*.
- **1.5.1.** 3% hydrogen peroxide for catalse test .
- **1.5.2.** Tetra-methyl-p-phenylene diamine 1% for oxidase test.
- **1.5.3**. Kovac's reagent: for the indole test.
- **1.5.4.** 0.04% methyl red solution: for methyl red test.
- 1.5.5. Solution I of 5.0% α -naphthol in absolute ethyl alcohol and solution II of 40% potassium hydroxide solution for Voges Proskauer test.
- **1.5.6**. 40% urea solution (Oxoid SR 20): for urease test.
- 1.5.7. Reagent I of 0.8% sulfanilic acid in 10 N acetic acid and reagent II of 0.5% α-naphthylamine in 5N acetic acid for nitrate reduction test.
- **1.5.8.** 10% and 40% bile salt.
- 1.5.9. 10% ferric chloride for phenyl alanine deaminase test.
- **1.5.10**. Streile paraffin oil for Arginine dehydrolase test.
- **1.5.11.** Phosophate buffer saline (*Emery et al., 1992*) for serogrouping of *Escherichia coli*.

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

- **1.6. Diagnostic** *Escherichia coli* **antisera:** The polyvalent and monovalent antisera used were:
 - Polyvalent 1: Contain: O1, O26, O86a, O111, O119, O127a, O128,
 - polyvalent 2: Contain: O₄₄, O₅₅, O₁₂₅, O₁₂₆, O₁₄₆, O₁₆₆,
 - polyvalent 3: Contain: O₁₈, O₁₁₄, O₁₄₂, O₁₅₁, O₁₅₇, O₁₂₈
 - polyvalent 4: Contain: O₆, O₂₇, O₇₈, O₁₄₈, O₁₅₉, O₁₆₈
 - Polyvalent 5: Contain: O₂₀, O₂₅, O₆₃, O₁₅₃, P₁₆₇
 - polyvalent 6: Contain: O₈, O₁₅, O₁₁₅, O₁₆₉
 - polyvalent 7: Contain: O_{28ac} , O_{112ac} , O_{124} , O_{136} , O_{144} and
 - polyvalent 8: Contain:O₂₉,O₁₄₃,O₁₅₂,O₁₆₄

It was obtained from Denka Seiken Co., LTD 3-4-2 Nihonbashikayaba – , Chuo. Ju, Tokyo, Japan.

1.7. Laboratory animals:

White mice:

Swiss Webster white mice weighting 18 - 22 grams were used to measure the virulence of bacterial isolates.

1.8. Diagnostic antibiotics and chemotherapeutic agents: (Oxoid):

Antimicrobial sensitivity discs were obtained from Oxoid. The sensitivity of bacterial isolates was tested against the following antimicrobial agent's disks: ciprofloxacin. $(3\mu g)$, cloxacillin $(5\mu g)$, chloramphenicol $(30\mu g)$, Doxycyline $(30\mu g)$, enrofloxacin $(5\mu g)$, flumequine $(30\mu g)$, Gentamicin $(10\mu g)$, norfloxacin $(10\mu g)$, penicillin $(10\mu g)$, streptomycin $(10\mu g)$ and trimethoprim $(5\mu g)$.

1.9. Sodium Dodecyl Sulphate Polyacrylamide Gel Electrophoresis (SDS-PAGE):

Determination of whole cell protein the molecular weight of the (*Klebsiella spp.* and *E. coli serogrouping*) by the SDS-PAGE according to (*Laemmli, 1970*).

i- Equipment:

1-Mini-protean 3 electrophoresis (Bio Rad).

2-Power supply.

ii-Reagents:

- 1- Acrylamide 30% stock solution : Stored in 4 °C. In brown bottle.
- 2- Separation buffer: PH adjusted to 8.8 with 6 N HCL, Stored in 4 °C. In brown bottle.
- 3- Stacking buffer: PH adjusted to 6.8 with 6 N HCL, stored in $4C^0$. in brown bottle.
- 4- Ammonium persulphate (APS): 400 μ g /ml (stored at- 20 C⁰ in small aliqutes).
- 5- TEMED (NNN'N' tetramethyl ethylene diamine).
- 6- Glycerol pure 100%.
- 7- Sample buffer (2x).

PH adjusted to 6.8, Stored at 4 C^0 in brown bottle.

- 8- Stock electrophoresis (10x).
- 9- Commassie blue stain: (0.25% commassie blue powder dissolved in destaining solution).

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

- 10- De-staining solution: 45% methanol. 50% glacial acetic acid and 50% distilled water.
- 11- Matrix contains 10% separating gel.

2. METHODS:

2.1. Collection and preparation of samples:

Samples were obtained from apparently healthy and diseased lungs which showing various macroscopic pathological changes pneumonia. The samples were collected separately in sterile plastic ice bags and transported immediately to the laboratory in ice box for bacteriological examination.

2.2. Bacteriological examination:

2.2.1. Cultivation:

The surface was sterilized by hot spatula then opened by sterile scalpel. Sterile platinum loop was inserted inside the opening then inserted into nutrient broth also one gm from sample was put in 10 ml of nutrient broth and incubated aerobically at 37°C for 24hrs. The suspected colonies were picked up and subculture into nutrient agar and specific media plates for each bacteria to confirm their purity then into nutrient agar slopes for identification.

2.2.2. Identification of bacterial isolates (Fineglod and Martin, 1982 and Quinn et al., 1994):

Cultivated plates were examined after 24-48 hrs incubation at 37°C for any bacterial growth. Colonies were identified according to:

2.2.2.1. Culture characters:

The cultural characteristic of each purified isolate were studied on solid madia according to their shape, size ,elevation ,structure of surface, colour, pigment and type of haemolysis .

2.2.2.2. Microscopical examination:

The isolated organisms were classified according to shape, size, arrangement and gram staining reaction into: Gram positive cocci, Gram positive non spore forming rods and Gram negative rods.

2.2.2.3. Biochemical identification:

2.2.2.3.3. Identification of Gram negative rods (Quinn et al., 2002):

2.2.2.3.3.1. Identification of *Pseudomonas species*.

Pyocyanine production (pigment production), Oxidase production, Gelatin liquefaction, Growth at 4°C and 41°C, Catalase test, Reduction of nitrate, Decarboxylation of arginine (Arginine dihydrolase) and sugar utilization test.

2.2.2.3.3.2. Differentiation of members of Family Enterobacteriaceae:

Indole production, Methyl red, Voges proskauer, Citrate utilization, Urease activity, Hydrogen sulphide production, Motility and swarming on blood agar, Gelatin liquefaction. and Sugar fermentation test.

2.2.2.3.3.3. Differentiation of *Pasteurella* spp. (*Barrow and Feltham*, 1993):

Motility, Catalse test, Oxidase test, Growth on the MacConkey agar plates, Sugar fermentation test using arabinose, lactose, maltose, mannitol, raffinose, salicin, sorbitol, sucrose and trehalose, Aesculin hydrolysis, Nitrate reduction test, Indole production test, Gelatin hydrolysis, Urease hydrolysis test and H_2S production by using lead acetate paper.

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

2.3. Serological identification:

2.3.1. Serogrouping of *Escherichia coli* isolates: (*Edwards and Ewing*, 1972).

All the suspected microorganisms were subjected to serological typing by slide agglutination test using standard polyvalent and monovalent *E. coli* antisera. Only fresh bacterial cultures 24hrs colonies onto nutrient agar medium were used and not from the selective medium.

Slide agglutination test:

When a colony gave a strongly positive agglutination with one of the pools of polyvalent serum, a further portion of it was inoculated into a nutrient agar slant and incubated to grow as a culture to test with monovalent sera.

2.4. Pathogenicity test of bacterial isolates in mice:

a) Preparation of bacterial suspension: (Stamp et al. 1959).

The bacterial suspension was made by plate washing technique. From the original culture, this was plated onto 10% sheep blood agar plate and then incubated for 24 hrs at 37°C. The inoculated plate was flooded with 5 ml saline and the colonies were removed from the solid medium by gentle rubbing with a glass rod, the resultant suspension were adjusted to be contained on average of 3 x 10^8 colony forming unite (CFU) per ml (adjusted to match a McFarland standard tube No.1).

b) Route of injection (Wessman, 1964):

Five mice were used for each isolate; they were injected from the prepared solution using sterile disposable plastic syringes. Each mouse was given 0.1 ml of bacterial suspension I/P in the group for the detection of pathogencity. The mortalities rates and the pathogencity rate were calculated.

2.5. Susceptibility of the isolated strains to various antimicrobial agents:

The disk diffusion technique was adapted according to *Fineglod* and Martin (1982).

2.5.1. Preparation of the standardized inoculums:

- Four or five typical colonies of similar morphology were transferred using a sterile loop to a tube containing 5 ml of sterile Mueller Hinton broth.
- The broth was incubated at 37°C for 2 8 hrs for its turbidity exceeds that of the standard McFerland tube 0.5 barium sulphate tube (0.5 ml of 1.175% barium chloride hydrate to 99.5ml of 1% sulphuric acid). The turbidity was adjusted to match a McFerland 0.5 barium sulphate standard tube by adding sterile saline using adequate light

2.5.2. Inoculation of the tested plates:

- A sterile cotton swab was dipped into the standardized bacterial suspension.
- Excess fluid was removed by rotating the swab with firm pressure against the inside wall of the tube above the fluid level.
- The swab was then used to streak the dried surface of Muller-Hinton agar plate in three different planes by rotating the plate approximately (60) degree each time to ensure an even distribution of the inoculation.
- The plate lids were placed and the inoculated plates were allowed to remain on a flat level surface undistributed for 3 to 5 minutes to allow the adsorption of excess moisture then the disks were applied.

2.5.3. Application of chemotherapeutic discs:

- With fine-pointed forceps, the selected antimicrobial discs were placed onto the inoculated plates.
- The discs were distributed evenly in a manner such as to be no closer than 15 mm from the edge of Petri dish and so that no two discs were closer than 24 mm from center to center. The plates were then incubated at 37°C for 24 hrs.

2.5.4. Interpretation of the zone of inhibition:

By measuring clear zone of inhibition of growth produced by diffusion from the discs into the surrounding medium.according to the *Koneman et al. (1983)* and *Oxoid Manual (1982)* Table (2).

Antibiotics and chemo	Symbol	Cono in ug	Diameter of inhibition zone					
therapeutic agents	Symbol	Conc. in µg	Sensitive	Resistance				
Chloramphenicol	С	30	≥ 18	≤ 12				
Ciprofloxacin	Cip	5	≥ 21	≤ 15				
Cloxacillin	OB	5	≥ 18	≤ 14				
Doxycycline	Do	30	19 – 23	15 - 18				
Enrofloxacin	Enr	5	≥ 18	≤ 14				
Flumequine	UB	30	≥16	≤ 13				
Gentamicin	CN	10	≥ 15	≤ 12				
Norfloxacin	Nor	10	≥ 17	≤ 12				
Pencillin G	Р	10	≥ 19	≤ 14				
Streptomycin	S	10	≥ 15	≤11				
Trimethoprim	Tr	5	≥ 16	≤ 10				

Table (2): Interpretation of zones of inhibition for antibacterial susceptibility.

All obtained from Oxoid Company.

2.6. Sodium Dodecyl Sulphate Polyacrylamide Gel Electrophoresis (SDS-PAGE):

It was performed according to the method described by *Laemmli* (1970).

2.6.1. Casting of the gel:

- 1. Fifteen ml of 12 % gel was prepared and the polymerization process occurred as soon as temed was added then poured in between the two glass plates carefully to avoid air bubbles.
- 2. Leaving a sufficient space for stacking gel.
- 3. After polymerization was completed (30 minutes.), the top of the gel was washed by deionized water for several times to remove non polymerized solution then removal of any water by using towel paper.
- 4. The stacking gel was prepared and then poured directly on the surface of polymerized separating gel.
- 5. A clean Teflon comp was immediately inserted into the stacking gel solution careful to avoid trapping any air bubbles. The gel was kept in a vertical position at room temperature till polymerization.

2.6.2. Preparation of the samples:

The samples 10 μ l / well was treated with the reducing buffer in ratio of 1:2 the treated samples were immersed in a boiling water for 2 minutes to insure protein denaturation.

2.6.3. Samples loading:

- 1. After polymerization of stacking gel was completed the Teflon comp was carefully removed and the formed wells washed by deionized water to remove any of the non polymerized solution.
- 2. The gel was mounted in the tank of electrophoresis apparatus. The running buffer was added to the top and bottom reservoirs and removal of any air bubbles was assured.
- 3. About, 10 μ l of each sample and marker were loaded with its loading buffer into wells of the gel. The power supply was turned on and the current was adjusted at 2 minutes Amp/sample and the voltage was adjusted at 100 volt.

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

4. The power supply was turned off when the blue line descends to the bottom of the rubber of the electrophoresis apparatus.

2.6.4. Commassie brilliant blue (R-250) 0.25: (Laemmli, 1970).

The gel was removed between the two glass blades carefully and placed in staining (Commassie brilliant blue) followed by gentle shaking (20 minutes.) and washed gently with deionized water.

The gel removed from deionized water and placed in destained solution (45% methanol, 50% glacial acetic acid and 50% distilled water) for 30-45 minutes.

Destaining was carried out several times until bands appeared and the gel cleared.

RESULTS

3.1. Incidences and types of positive cases yielded single or mixed bacterial isolates from lungs of slaughtered camels:

Table (3): The incidences and types of positive cases yielded single or mixed bacterial isolates from lungs of slaughtered camels.

Camel health status	No. of examined samples	No. of positive samples	º⁄₀*	No. of samples yielded single isolate	%**	No. of samples yielded mixed bacterial isolate	%**	No. of isolates	Gram positive bacterial isolates No.	%***	Gram negative bacterial isolates No.	0⁄0***
Apparently healthy	205	124	60.49	74	59.68	50	40.32	174	80	45.98	94	54.02
Diseased (pneumonic)	103	79	76.70	30	37.97	49	62.03	128	52	40.63	76	59.38
Total	308	203	65.91	104	51.23	99	48.77	302	132	43.71	170	56.29

* The percentage was calculated according to number of examined samples of each camel health status.

** Percentage was calculated according to total No. of positive samples of each camel health status.

*** Percentage was calculated according total number of isolates (174) from apparently healthy and (128) from pneumonic camel lungs.

 Table (4): The incidences and types of bacterial isolates recovered from apparently healthy and pneumonic lungs of slaughtered camels.

Types of bacteria	Apparently healthy lur	ngs (124 samples)	Pneumonic lungs (79 samples)										
Types of bacteria	No. of isolates	%	No. of isolates	%										
b) Gram negative isolates														
Citrobacter freundii	11	6.32	5	3.91										
Citrobacter spp.	8	4.6	3	2.34										
E. coli	29	16.67	20	15.63										
Enterobacter cloacae	7	4.02	3	2.34										
Klebsiella oxytoca	8	4.60	6	4.69										
Klebsiella ozaenae	3	1.72	2	1.56										
Klebsiella pneumoniae	0	0	12	9.38										
Pseudomonas aeruginosa	6	3.45	10	7.81										
Mannheimia haemolytica	0	0	2	1.56										
Pasteurella multocida	0	0	1	0.78										
Proteus penneri	9	5.17	4	3.13										
Proteus vulgaris	5	2.87	2	1.56										
Yersinia enterocolitica	8	4.6	6	4.69										
Total	94	54.02	76	59.38										

N.B: Total number of bacterial isolates from apparently healthy lung were 171 including gram positive &gram negative ones while total number of bacterial isolates from pneumonic lungs were 128 including also gram positive & and gram negative bacteria.

*The percentage was calculated according to the total number of bacterial isolates from each camel health status.

Table (5): The incidences and types of single isolated gram negative isolate recovered from both apparently healthy and pneumonic lungs of slaughtered camels.

Isolated microorganisms	Total posit from	ive samples lungs		mples from nealthy lungs	Positive samples from pneumonic lungs										
	No.	%*	No.	%	No.	%									
b) Gram negative isolates	b) Gram negative isolates														
Citrobacter freundii	7	6.73	7	9.46	0	0									
Citrobacter spp.	5	4.81	5	6.76	0	0									
E. coli	28	26.92	21	28.38	7	23.33									
Klebsiella oxytoca	3	2.89	3	4.05	0	0									
Klebsiella ozaenae	1	0.96	1	1.35	0	0									
Klebsiella pneumonia	6	5.77	0	0	6	20									
Mannheimia haemolytica	2	1.92	0	0	2	6.67									
Pasteurella multocida	1	0.96	0	0	1	3.33									
Pseudomonas aeruginosa	9	8.65	4	5.41	5	16.67									
Yersinia enterocolitica	2	1.92	2	2.7	0	0									
Total	104	100	74	71.15	30	28.85									

* The percentage was calculated according to the total number of positive samples(104) in all lung samples, (74) in apparently healthy lungs and (30) in pueumonic lungs.

Table (6): The incidences and types of mixed bacterial isolates recovered from

 both apparently healthy and pneumonic lungs of slaughtered camels.

		oositive rom lungs		mples from	positive samples from pneumonic lungs				
Isolated microorganisms				nealthy lungs	-)			
	No.	%	No.	%	No.	%*			
A.rcanobacterium pyogenes + Citrobacter spp.	3	3.03	0	0	3	6.12			
C.itrobacter freundii + Yersinia. enterocolitica	9	9.1	4	8	5	10.2			
Diplococcus pneumoniae + E. coli	2	2.02	0	0	2	4.09			
Diplococcus. pneumoniae + Pseudomonas aeruginosa	4	4.04	0	0	4	8.16			
Klebsiella oxytoca + E. coli	3	3.03	0	0	3	6.12			
Klebsiella ozaenae + E. coli	1	1.01	0	0	1	2.04			
Klebsiella ozaenae + Y. enterocolitica	3	3.03	2	4	1	2.04			
Micrococcus spp.+ Citrobacter spp.	3	3.03	3	6	0	0			
Micrococcus spp. + E. coli	4	4.04	2	4	2	4.09			
Micrococcus spp.+K. pneumoniae	1	1.01	0	0	1	2.04			
Non hemolytic Streptococci + Proteus vulgaris	5	5.05	3	6	2	4.09			
Proteus penneri + Micrococcus spp.	13	13.13	9	18	4	8.16			
Pseudomonas aeruginosa + E. coli	1	1.01	0	0	1	2.04			
Pseudomonas aerugnosia + Proteus vulgaris	2	2.02	2	4	0	0			
Staphylococcus aureus + D. pneumoniae	2	2.02	0	0	2	4.09			
Staphylococcus aureus + E. coli	10	10.10	6	12	4	8.16			
Staphylococcus aureus + E. cloacae	10	10.10	7	14	3	6.12			
Staphylococcus aureus + K. oxytoca	8	8.08	5	10	3	6.12			
Staphylococcus aureus + K. pneumoniae	5	5.05	0	0	5	10.2			
Staphylococcus. aureus + non hemolytic Streptococci	10	10.10	7	14	3	6.12			
Total	99	100	50	50.5	49	49.5			

*The Percentage was calculated according to the total number of positive samples (99) in all lungs samples, (50) in apparently lungs and (49) in pneumonic lungs.

	Apparently healthy lungs		Pneumonic lungs						
Strain	No. of isolates	%	S <mark>t</mark> rain	No. of isolates	%*				
O ₁₁₁	6	20.68	O ₁₁₁	2	10				
O ₁₂₅	5	17.24	O ₁₂₅	3	15				
O ₁₆₆	5	17.24	O ₁₆₆	4	20				
O ₁₄₆	4	13.79	O ₁₄₆	2	10				
O ₅₅	3	10.35	O ₅₅	5	25				
O ₁₁₅	3	10.35	O ₁₁₅	1	5				
O ₁₁₉	3	10.35	O ₁₁₉	3	15				
Total	29	100	Total	20	100				

 Table (7): Serogrouping of *E. coli* isolates from apparently healthy and pneumonic lungs of camels.

*The Percentage was calculated according to number of isolates of each camel health status.

 Table (9): Pathogenicity test of Gram negative bacterial isolates recovered from lungs of slaughtered camels.

Examined bacteria]	No. of (died m	ice/ day	7		Total	%
Examineu bacteria	1	2	3	3 4		6	7	Total	70*
1.E. coli O ₅₅	3	2	0	0	0	0	0	5/5	100%
2.E. coli O ₁₁₅	2	1	2	0	0	0	0	5/5	100%
3.Klebsiella pneumoniae	0	2	1	1	1	0	0	5/5	100%
4.Mannheimia haemolytica	4	1	0	0	0	0	0	5/5	100%
5.Pasteurella multocida	5	0	0	0	0	0	0	5/5	100%
6.Pseudomonas aeruginosa	1	2	2	0	0	0	0	5/5	100%
7.E. coli O ₁₂₅	0	1	2	0	0	0	0	3/5	60%
8.E. coli O ₁₄₆	2	1	0	0	0	0	0	3/5	60%
9.E. coli O ₁₆₆	1	2	0	0	0	0	0	3/5	60%
10.Citrobacter freundii	0	0	1	1	0	0	0	2/5	40%
11.E. coli O ₁₁₁	1	0	0	0	1	0	0	2/5	40%
12.Enterobacter cloacae	0	0	1	1	0	0	0	2/5	40%
13.Klebsiella oxytoca	0	0	1	1	0	0	0	2/5	40%
14.Klebsiella ozaenae	0	0	1	1	0	0	0	2/5	40%
15.Yersinia enterocolitica	0	1	1	0	0	0	0	2/5	40%
16.E. coli O ₁₁₉	0	1	0	0	0	0	0	1/5	20%
17.Proteus vulgaris	0	0	1	0	0	0	0	1/5	20%
18. Citrobacter spp.	0	0	0	0	0	0	0	0	0%
19.Proteus penneri	0	0	0	0	0	0	0	0	0%

*The percentage was calculated according to the number of examined mice of each strain

Table (11): The antibiotic and chemotherapeutic agents of Gram negativebacterial isolates recovered from apparently healthy andpneumonic lungs of slaughtered camels.

Tested pathogens	ndii			E. coli						acae			е	a		ri	is	a	3a	
Antibiotic discs	Citrobacter freundii	Citrobacter sp.	O ₅₅	O ₁₁₁	O ₁₁₅	O ₁₁₉	O ₁₂₅	O ₁₄₆	O ₁₆₆	Enterobacter cloacae	K. oxytoca	K. ozaenae	K. pneumoniae	M. haemolytica	P. multocida	Proteus penneri	Proteus vulgaris	Ps. Aeruginosa	Y. enterocolitica	Total
Chloramphenicol	R	++	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	1
Ciproflocacin	+	R	R	R	R	+	R	+	++	++	R	R	R	R	R	+	++	R	R	7
Cloxacillin	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	0
Doxycycline	R	R	R	R	R	R	R	R	R	R	R	R	R	++	++	R	R	R	++	3
Enrofloxacin	+	++	++	+++	++	+++	++	++	+++	++	+++	++	++	+++	+	++	++	++	+	19
Flumequine	R	R	R	R	R	R	+	R	R	+	++	+++	+++	R	R	R	R	R	R	5
Gentamicin	R	R	++	+	+	++	R	+	++	R	+	+	R	R	R	R	R	R	R	8
Norfloxacin	R	++	+	+	++	++	+	++	++	++	++	++	+++	+	R	+	++	R	R	15
Pencillin	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	0
Streptomycin	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	0
Trimethoprim	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	0

-R= Resistant -+= Sensitive -++= Wide Sensitive zone -+++= Very wide Sensitive zone

Ten strains including 3 Klebsiellae :*Klebsiella pneumoniae*, *Klebsiella oxytoca*, *Klebsiella ozaenae* and 7 strains of *E. coli* including : *E. coli* O55, *E. coli* O111, *E. coli* O115, *E. coli* O119, *E. coli* O125, *E. coli* O146 and *E. coli* O166 isolated from camel lung samples.

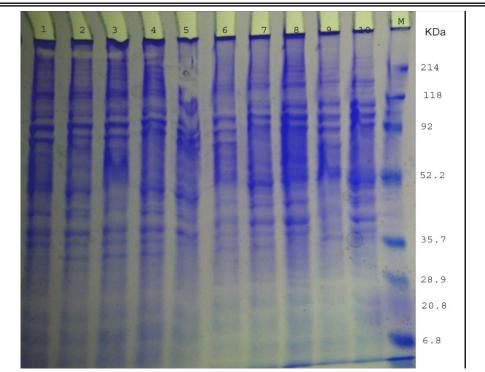


Photo (1): whole cell protein mixture were run on poly acrylamide gel stained with Commassie brilliant blue for detection of whole cell protein profiles lane 1 *E. coli* O55, lane 2 *E. coli* O111, lane 3 *E. coli* O115, lane 4 *E. coli* O119, lane 5 *E. coli* O125, lane 6 *E. coli* O146, lane 7 *E. coli* O166, lane 8 *Klebsiella oxytoca*, lane 9 *Klebsiella pneumoniae* and lane 10 *Klebsiella ozaenae* and lane 11, molecular weight marker Bio-Rad (214-118- 92- 52.2- 35.7- 28.9- 20.8- 6.8 KDa).

DISCUSSION

Bacterial infection of the lungs is one of the main causes of pneumonia in camels (*Rana et al., 1993; Thabet, 1994; Alhendi, 2000* and *Seddek, 2002*).

The present study is a trial to find the pathogenic bacteria that are associated with camel's lungs, also to study the pathogenicity and antibacterial sensitivity of the bacterial isolates to control bacterial pneumonia in camels.

Several hygienic factors may be predisposing causes of respiratory disease. These including long period without food, keeping in dirty stalls *Baba et al.*, *1994*) and transportation as camels come yearly from North of Sudan to south of Egypt and collected then remained for several weeks in daraw quarantine, Aswan, Egypt . Cold, raining and dust during winter are predisposing factors for respiratory infection (*Ning et al. 1998*).

As clear from **Table (5)** that bacteriological examination of 308 camel lung samples (205 apparently healthy and 103 pneumonic) revealed that 203 samples were bacteriologically positive with an incidence of 65.91%. These result was similar to that reported by *Mahmoud et al. (1988)* who found that 67.3% of lung samples were bacteriologically positive but Lower incidences were recorded by *Ebtesam et al. (2009)* and *Moustafa (2004)* who obtained 47.06% positive samples , while higher prevalence was recognized by *Amany (2000)* who had 86.24% positive samples.

Among 203 bacteriological positive lung samples, 124 (60.49%) of apparently healthy lungs were bacteriological positive which partially confined with *Thabet (1994)* who his incidence was (71.8%) and disagreed with *Azzam and Zaki (2006)* who recorded lower prevelance (26.66%).

Meanwhile,79 (76.7%) of pneumonic lungs were culturally positive for bacteria which partially agreed with *Azzam and Zaki (2006)* who found that (79%) of pneumonic samples were bacteriological positive. Higher incidences were estimated by *Moustafa (2004)* who recorded (96%) of pneumonic samples were bacteriological positive. Lower percentages were recorded by *Fatma et al. (2001)* who found (56%) of pneumonic samples were bacteriological positive.

The isolated bacteria may be present singly or mixed with other. It was noticed from the same table that out of 203 bacterial positive lungs, 104 (51.23%) samples had single infection that partially agreed with *Amany (2000)* who recovered single isolates from (65%) of lung samples. On the other hand, 99 (48.77%) samples yielded mixed infection who found that was nearly similar to *Nagi et al. (1997)* who found that (43.85%) yielded mixed infected samples.

In the present study, the incidences of samples yielded single and mixed infection from apparently healthy lungs of camels were 74 (59.68%) and 50 (40.32%), respectively, this result disagreed with *Azzam and Zaki (2006)* who recorded that the incidence were (85.71%) and (14.29%) for single and mixed samples of apparently healthy lung of camels, respectively.

On the other hand, prevalence of pneumonic lung samples yielded single isolates was 30 (37.97%) which was approximately similar to *Moustafa* (2004) who recorded the incidence of samples yielded single isolates from pneumonic lungs was (41.7%).

The incidence of samples yielded mixed isolates from pneumonic lungs was 49 (62.03%) which confined to some extent *Moustafa* (2004) who found that (58.3%) of samples yielded mixed isolates from pneumonic lungs.

These findings declared that the highest samples yielded single bacterial isolates than mixed ones were found in apparently healthy lungs, which agreed with *Thabet (1994)*. Meanwhile, the contrast occurs as the highest samples yielded mixed isolates were more than that yielded single

The current investigations revealed that 302 bacterial isolates recovered from lungs of slaughtered camels classified as 132 (43.71%) Gram positive and 170 (56.29%) Gram negative isolates out of which, 174 were recovered from apparently healthy lungs that classified as 80 (45.98%) of Gram positive and 94 (54.02%) of Gram negative isolates. These results were partially confined with *Moustafa (2004)* and disagreed with *Amany (2000)* who isolated (78.13%) of Gram positive and (21.88%) of Gram negative bacteria.

However, the number of isolates were 128 from pneumonic lungs, distributed as 52 (40.63%) of Gram positive and 76 (59.38%) of Gram negative isolates which were nearly in accordance with *Amer et al.* (2002) who found that the incidence of Gram positive were (54.7%) and (45.33%) of Gram negative isolates respectively, and disagreed with *Al-Doughaym et al.* (1999) who had (82.4%) of Gram positive and (15%) of Gram negative isolates from pneumonic lungs.

These variations in our data compared with others may be due to location, country, husbandary, feeding and immunity.

Meanwhile, the highest number of Gram negative bacteria obtained from apparently healthy lungs was *E. coli* with an incidence of 29 (16.67%) which partially agreed with *Aly et al.* (2004) who found that *E. coli* was the main Gram negative isolate from apparently healthy lungs with an incidence of (22.2%) and disagree with *Moustafa* (2004) who isolated *E. coli* with a percentage of (2.94%) from apparently healthy lungs.

Many other Gram negative isolates were recovered from apparently healthy lungs such as *Citrobacter freundii* with an incidence of 11 (6.32%), *Proteus penneri* 9 (5.17%), *Citrobacter spp., K. oxytoca* and *Y. enterocolitica* with an incidence of 8(4.6%) in each, *Enterobacter cloacae* 7 (4.62%), *Pseudomonas aeruginosa* 6 (3.45%), *P. vulgaris* 5 (2.87%) and finally, *K. ozaenae* 3 (1.72%).

Isolation of *Citrobacter spp.* from apparently healthy lungs of camels was previously recognized by *Amany (2000)*. *Pseudomonas aeruginosa* was isolated by *Ebtesam et al. (2009)*.

P. vulgaris from apparently healthy lungs of camels was previously recorded by *Moustafa* (2004) but with different incidences.

Klebsiella spp. were isolated from apparently healthy lungs of camels by *Hanea and Omnia (2002)* and *Thabet (1994)* who found *Klebsiella spp.* in incidence of (4.9%).

New isolates were isolated from apparently healthy camel lungs such as *P. penneri*, *K.oxytoca*, *K. ozaenae*, *Entrerobacter cloacae* and *Y. enterocolitica* that were not previously recorded by other authors.

These results indicated that apparently healthy lungs act as a reservoir for many species of pathogenic and potential pathogenic microorganisms that under stress factors such as changes in the hygiene, environmental and climatic conditions may play a role in the onset of pneumonia **Zubair et al. 2004**). This concept was supported in the present study by the fact that a number of bacteria were isolated from 124 samples which showed no pathological lesions.

On the other hand, the highest number of Gram negative isolates was *E. coli* with an incidence of (15.63%) that agreed with *Amer et al.* (2002) who found that *E. coli* was the predominant isolate of Gram negative bacteria recovered from pneumonic lungs but with different percentages.

The other Gram negative bacteria incriminated in camel pneumonia were *K. pneumoniae* 12 (9.38%), *Ps. aeruginosa* 10 (7.81%), *K. oxytoca* and *Y. enterocolitica* with an incidence of 6 (4.69%) in each, *Citrobacter freundii* 5 (3.91%), *P. penneri* 4 (3.13%), *Citrobacter spp.* and *Enterobacter cloacae* with an incidence of 3 (2.34%) in each, *K. ozaenae* and finally *P. multocida* with an incidence of 1 (0.78%) in each.

Incidence of *E. coli* was partially agreed with *Seddek (2002)* who isolated *E. coli* from 10.58% of pneumonic samples while disagreed with *Al-Doughaym et al. (1999)* who detected *E. coli* with an incidence of 6.2%.

Klebsiella pneumoniae was observed by *Nagi* (2004). Similar incidence was showed by *Al-Doughaym* (1999) who obtaine *Klebsiella pneumoniae* from (10.9%) of pneumonic samples and disagreement occurred with *Amer et al.* (2002) who obtained *Klebsiella pneumoniae* from (3.8%) of pneumonic lungs.

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

Pseudomonas aeruginosa was detected from pneumonic lungs by *Al-Tarazi and El-Sheikh (2006)*. The incidence of *Pseudomonas aeruginosa* is partially agreed with *Seddek (2002)* who obtained *Pseudomonas aeruginosa* (8.82%) from pneumonic lungs, while disagreed with *Asil et al. (2004)* who found *Pseudomonas aeruginosa* in an incidence of (1%) of pneumonic lungs.

Klebsiella oxytoca and *Klebsiella ozaenae* were previously recovered by *Al-Tarazi* (2001) from pneumonic lungs.

Lower percentage of *Yersinia enterocolitica* was recorded by *Fatma et al. (2001)* who isolated *Yersinia enterocolitica* from (1.26%) in pneumonic lung samples whereas higher incidence (9.4%) was recorded by *Amer et al. (2002)*.

Citrobacter spp. were observed by *Abubakr et al. (1999)*. The incidence was agreed with *Seddek (2002)* who isolated Citrobacter spp. from (2.35%) of pneumonic samples and partially coincided with *Fatma et al. (2001)* who revealed it in a lower incidence of (1.26%).

Citrobacter freundii was showed by Wernery and Kaaden (2002). The incidence was partially agreed with Fatma et al. (2001) who found Citrobacter freundii in an incidence of 2.53% from pneumonic lungs of camels.

Proteus spp. were previously observed from pneumonic lungs by *Kane et al. (2005)*.

Proteus vulgaris was recognized by *Mahmoud et al. (1988)* while Proteus penneri was previously obtained by *Fatma et al. (2001)* with lower percentage (1.26%). The incidence of *Proteus vulgaris* is partially

agreed with *Fatma et al.* (2001) who isolated it with an incidence of (1.26%) while higher percentage (9.46%) was isolated by *Zaitoun* (1986).

Pasteurella spp. were isolated by many searchers from pneumonic lungs such as *Asil et al.* (2004) and *Nagi* (2004). Also, *Pasteurella multocida* was recognized by *Al-Ani* (1990) whilst *Mannheimia haemolytica* was detected by *Al-Tarazi and El-Sheikh* (2006).

In the present work, *Mannheimia haemolytica* was recovered with a percentage of 1.56% which partially agreed with *Fatma et al. (2001)* who isolated it in incidence of (1.17%) from pneumonic lung samples.

Meanwhile, *Pasteurella multocida* was isolated with a percentage of 0.78% which is lower than that recorded by *Seddek (2002)* and *Susan et al. (2000)* who isolated (14.2%) from pneumonic lungs. The low percentage of isolated Pasteurella spp. was supported by *Magda (1998)* who stated that camels are less susceptible to Pasteurella infection than other animals.

Enterobacter spp. were revealed by *Asil et al. (2004)* from pneumonic lungs with various prevalences.

It was clear from **Table (5)** that 74 samples yielded single bacteria l isolates from apparently healthy lungs of slaughtered camels that were classified into two categories, Gram positive and Gram negative bacterial isolates.

On the other hand, the highest incidence of Gram negative single bacterial isolates was *E. coli* 21 (28.38%) followed by *Citrobacter freundii* 7 (9.46%), Citrobacter spp. 5 (6.76%), *Pseudomonas aeruginosa* 4 (5.41%), *Klebsiella oxytoca* 3 (4.05%), *Yersinia enterocolitica* 2 (2.7%) and *Klebsiella ozaenae* 1 (1.35%).

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

On the other hand, the most prevalent single isolate of Gram negative bacteria from pneumonic lungs of camels was *E. coli* in 7 samples with an incidence of (23.33%) then *Klebsiella pneumoniae* 6 (20%), *Pseudomonas aeruginosa* 5 (16.67%), *Mannheimia. haemolytica* 2 (6.67%) and *Pasteurella multocida* 1 (3.33%).

These findings were partially agreed with *Azzam and Zaki (2006)* who isolated *E. coli*, *Pasteurella haemolytica* and *Pasteurella multocida* with incidence of 26%, 10% and 5%, respectively.

Lower incidences of *Pseudomonas aeruginosa* and *Klebsiella pneumoniae* were showed by *Moustafa* (2004), *Azzam and Zaki* (2006) and *Amany* (2000) who noticed *Pseudomonas aeruginosa* and *Klebsiella pneumoniae* as single isolates in pneumonic samples with percentages of 3.85% and 1.92%, respectively.

Higher incidences of *Pasteurella multocida* obtained by *Amany* (2000), *Selim* (2003) and *Moustafa* (2004) who isolated *Pasteurella multocida* from (8.3%) of pneumonic samples.

However, **Table (5)** noticed that out of 104 samples revealed single isolates of bacteria, 74 samples from apparently healthy lungs with an incidence of (71.15%) and 30 samples from pneumonic lungs with an incidence of (28.85%) yielded single isolates. The higher number of samples yielded single isolates was recovered from apparently healthy lungs. The recovered organisms were classified into two categories, Gram positive and Gram negative bacterial isolates. The most common Gram negative single isolates were *E. coli* in 28 lung samples with an incidence of 26.92% followed by *Pseudomonas aeruginosa* 9 (8.65%), *Citrobacter freundii* 7 (6.73%), *Klebsiella pneumoniae* 6 (5.77%), Citrobacter spp. 5

(4.81%), *Klebsiella oxytoca* 3 (2.89%), *Mannheimia haemolytica* and *Yersinia enterocolitica* with an incidence of 2 (1.92%) in each and finally *Klebsiella ozaenae* and *Pasteurella multocida* with an incidence of (0.96%) in each.

It was declared from **Table** (6) that bacteria isolated from apparently healthy lungs may occur in a mixed form mainly as a combination between *Proteus penneri* and Micrococcus spp. (18%), *Staph. aureus* with each of *Enterobacter cloacae* (14%), non haemolytic streptococci (14%), *E. coli* (12%) and *Klebsiella oxytoca* (10%) or another combination between Micrococcus spp. with each of Citrobacter spp. (6%) and *E. coli* (4%). Also, there was a combination between *Proteus vulgaris* with each of non haemolytic streptococci (6%) and *Pseudomonas aeruginosa* (4%) as well as *Yersinia enterocolitica* with each of *Citrobacter freundii* (8%) and *Klebsiella ozaenae* (4%).

The current investigation from **Table (6)** showed that the bacteria causing pneumonia in camel may occur in a mixed form consisted mainly of *Citrobacter freundii* with *Yersinia enterocolitica* or *Staphylococcus aureus* with *Klebsiella pneumoniae* with an incidence of (10.2%) in each.

Also, there were combinations between *Staph. aureus* with each of *E. coli* (8.16%), *Enterobacter cloacae* (6.12%), *Klebsiella oxytoca* (6.12%), non haemolytic streptococci (6.12%) and *Diplococcus pneumoniae* (4.09%).There was sharing between Micrococcus spp. and each of *Proteus penneri* (8.16%), *E. coli* (4.09%) and *Klebsiella pneumoniae* (2.04%) as well as *Diplococcus pneumoniae* with each of *Pseudomonas aeruginosa* (8.16%) and *E. coli* (4.09%).

The incidences of *Diplococcus pneumoniae* with *Pseudomonas aeruginosa* and *Staphylococcus aureus* with each of *E. coli* and *Enterobacter cloacae* were higher than those isolated by *Fatma et al.* (2001) and *Seddek* (2002) who recognized lower percentage of (1.44%), (2.89%) and (1.44%), respectively.

Combination between *Diplococcus pneumoniae* with *E. coli* was agreed with that recorded by *Azzam and Zaki (2006)* who had a percentage of (4%) from pneumonic samples.

Combination between *Staph. aureus* with non haemolytic Streptococci was previously recognized by *Fatma et al. (2001)* but with lower percentage (1.67%).

In the present study, it was noticed that *Staphylococcus aureus* was the highest mixed isolate isolated from pneumonic lungs of camels which is supported by *El-Haenaeey et al. (1994)*.

Amany (2000), Fatma et al. (2001), Seddek (2002), Selim (2003), Moustafa (2004) and Azzam and Zaki (2006) isolated different mixed bacterial isolates from pneumonic lungs including Staph. aureus with Pasteurella multocida, E. coli with Klebsiella pneumoniae, Strept. pneumoniae with Coryn. pyogenes, Micrococcus spp. with Citrobacter freundii and Staphylococcus aureus with Yersinia enterocolitica. This may be attributed to changes in environment, hygienic conditions, nutritional factors and immune status of the animal (Fatma et al. 2001).

It was clear from **Table (6)** that mixed isolates recovered from both apparently healthy and pneumonic lungs were mainly composed of *Proteus penneri* and Micrococcus spp. with an incidence of 13 (13.13%) followed by *Staphylococcus aureus* with each of *E. coli, Enterobacter*

cloacae and non haemolytic Streptococci with an incidence of 10 (10.10%) for each, Citrobacter freundii with Yersinia enterocolitica 9 (9.1%), Staph. aureus with Klebsiella oxytoca 8 (8.08%), non haemolytic streptococci with Proteus vulgaris 5 (5.05%), Staph. aureus with Klebsiella pneumoniae 5 (5.05), Diplococcus pneumoniae with Pseudomonas aeruginosa 4 (4.04%) and Micrococcus spp with E. coli 4 (4.04%) while, Arcanobacterium pyogenes with Citrobacter spp., Klebsiella oxytoca with E. coli, Klebsiella ozaenae with Yersinia enterocolitica and Micrococcus spp. with Citrobacter spp. were equal in an incidence of 3 (3.03%) in each. Also, the prevalence of Diplococcus pneumoniae with E. coli was equal to that of Pseudomonas aeruginosa with Proteus vulgaris and that of Staph. aureus with Diplococcus pneumoniae in two samples with an incidence of 2 (2.02%) in each.

The least mixed isolates were *Klebsiella ozaenae* with *E. coli*, *Micrococcus* spp. with *Klebsiella pneumoniae* and *Pseudomonas aeruginosa* with *E. coli* in one sample with an incidence of 1(1.01%)in each.

It was clear from **Table (7)** that serological identification of all the strains of *E. coli* isolated from apparently healthy and pneumonic lungs of slaughtered camels revealed 29 isolates from apparently healthy and 20 from pneumonic lungs.

There variations in incidence of serogroups isolated from apparently healthy lungs as *E. coli* O_{111} was the most predominant serogroup with an incidence of 6 (20.68%) followed by *E. coli* O_{125} and *E. coli* O_{166} with an incidence of 5 (17.24%) for each, *E. coli* O_{146} with an incidence of 4 (13.79%) then *E. coli* O_{55} , *E. coli* O_{115} and *E. coli* O_{119} with an incidence of 3 (10.35%) in each.

Meanwhile, in pneumonic lungs, the most predominant serogroup was *E. coli* O_{55} with an incidence of 5 (25%) followed by *E. coli* O_{166} with an incidence of 4 (20%), *E. coli* O_{119} and *E. coli* O_{125} with an incidence of 3 (15%) in each, *E. coli* O_{111} and *E. coli* O_{146} with an incidence of 2 (10%) in each and finally, *E. coli* O_{115} with an incidence of (5%).

This assumption was supported by *El-Battrawy et al.* (1992) who isolated *E. coli* O ₁₁₁ and *E. coli* O₁₂₅ from healthy and diseased animals and partially agreed with *Fatma et al.* (2001) in isolation of *E. coli* O₁₂₅ and *E. coli* O₁₆₆ and disaggreed with *Thabet* (1994), *Amany* (2000) and *Azzam and Zaki* (2006) who isolated *E. coli* O₈₆ and *E. coli* O₁₂₆ from pneumonic lungs of camels.

Regarding to the pathogenicity of the bacterial isolates, it was clear from **Table (8 and 9)** that the highest pathogenic bacteria of Gram positive bacterial isolates were *Arcanobacterium pyogenes* and *Diplococcus pneumoniae* with an incidence of (100%) in each, followed by *Micrococcus* spp. (60%), non haemolytic streptococci and *Staphylococcus aureus* with an incidence of (40%) in each.

Meanwhile, the highest pathogenic bacteria of Gram negative bacterial isolates were *E. coli* O_{55} , *E. coli* O_{115} , *Klebsiella pneumoniae*, *Mannheimia haemolytica*, *Pasteurella multocida* and *Pseudomonas aeruginosa* with an incidence of (100%) for each followed by *E. coli* O_{125} , *E. coli* O_{146} and *E. coli* O_{166} with an incidence of (60%) in each, *Citrobacter freundii*, *E.coli* O_{111} , *Enterobacter cloacae*, *Klebsiella oxytoca*, *Klebsiella ozaenae* and *Yersinia enterocolitica* with an incidence of (40%) in each as well as *E. coli* O_{119} and *Proteus vulgaris* with an incidence of 20% in each.

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

On the other hand, *Citrobacter spp.* and *Proteus penneri* were not being found to cause deaths among mice.

These findings agreed with *Buchnev et al. (1987), Refai (1990), Amany (2000), Omnia and Amer (2003)* and *Moustafa (2004)* who revealed death of (100%) of injected mice by *Pasteurella multocida* and partially agreed with *Susan et al. (2000)* who found that 81% of *Pasteurella multocida* and 33% of *Pasteurella haemolytica* had killed (100%) of mice within 24 hours post injection apparently healthy and diseased camels.

Concerning in-vitro antibacterial sensitivity, it was clear from **Table (10 and 11)**, that there is a marked difference between the sensitivity to antibiotics between different bacterial isolated from apparently healthy and pneumonic lungs of slaughtered camels.

These findings were concided with *Amany (2000)*, *Al-Tarazi and El-Sheikh (2006)* who found that most of isolates from pneumonic lungs of camels were sensitive to Ciprofloxacin, Enrofoxacin and Gentamicin and partially agreed with *Hanea and Omnia (2002)*.

It was noticed from **Table (13)** that the most effective antibiotics for Gram negative isolates from apparently healthy and pneumonic lungs of slaughtered camels were Enrofloxacin, Norfloxacin then Gentamicin, Ciprofloxacin and Flumequine then Doxycycline and Chloamphenicol but most of bacterial isolates were resistant to Cloxacillin, Penicillin, Streptomycin and Trimethoprim.

These findings were coincided with *Amer et al.* (2002) and *Hawari* (2008).

Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

E. coli serogroups were highly sensitive to Enrofloxacin, Norfloxacin also; Gentamicin except E. coli O₁₂₅, while E. coli O₁₁₉, E. coli O₁₄₆ and E. coli O₁₆₆ were sensitive to Ciprofloxacin and E. coli O₁₂₅ was sensitive to Flumequine.

Similar observations were noticed with Moustafa (2004) who found that *E. coli* was sensitive to Enrofloxacin and Gentamicin.

Enterobacter cloacae isolates were sensitive to Ciprofloxacin, Enrofloxacin and Norfloxacin followed by Flumequine while were resistant to Chloramphenicol, Cloxacillin, Doxycycline, Gentamicin, Penicillin, Streptomycin and Trimethoprim.

As shown from this table that the most effective antimicrobial agents against Klebsiella oxytoca, Klebsiella ozaenae and Klebsiella .pneumoniae were Enrofloxacin, Flumequine and Norfloxacin. K.oxytoca and K.ozaenae were sensitive to Gentamicin while K. pneumoniae was resistant. They were resistant to the other antimicrobial agents.

Hanea and Omnia (2002) who found that Klebsiella spp. were sensitive to Enrofloxacin, Gentamicin and Penicillin.

Klebsiella pneumoniae was sensitive to Enrofloxacin, Flumequine and Norfloxacin and resistant to Gentamicin which agreed with Abd El-Motelib and El-Zanaty (1993).

Mannheimia haemolytica was sensitive to Doxycycline and Enrofloxacin and Norfloxacin and resistant to the other antimicrobial agents.

Pasteurella *multocida* was sensitive Doxycycline to and Enrofloxacin and resistant to other antimicrobial agents. This agreed with Ahmed et al. (1986) who found that it was resistant to Penicillin and Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

partially agreed with *Moustafa* (2004) who found that *Pasteurella multocida* was sensitive to Chloramphenicol and Gentamicin and resistant to Streptomycin while disagreed with *Selim et al.* (2003) and *Azzam and Zaki* (2006) who revealed that *Pasteurella multocida* was sensitive to Norfloxacin.

Proteus penneri and *Proteus vulgaris* were sensitive to Ciprofloxacin, Enrofloxacin and Norfloxacin and resistant to the other types of antimicrobial agents.

Pseudomonas aeruginosa isolates were sensitive only to Enrofloxacin and resistant to the other types of antimicrobial agents which supported by *Thabet (1994)* and *Amer et al. (2002)* who found that *Pseudomonas aeruginosa* was sensitive to Enrofloxacin.

Yersinia enterocolitica was highly sensitive to Doxycyline and Enrofloxacin, Meanwhile, It was resistant to the other types of antimicrobial agents which partially agreed with *Amer et al.* (2002).

It was clear that most of bacterial isolates were resistant to most antibiotics and this may be attributed to wrong dosage, duration of treatment and route of administration (*Amstutz et al., 1982*).

These variations in sensitivity may be due to bacterial nature, distribution of bacteria or regional difference and plasmid helping in the formation of resistant strains. In consideration of the most effective in vitro antibacterial agents against all tested isolates, Enrofloxacin should be the drug of choice in controlling pneumonia in camel.

In this work, the whole cell protein of ten strains including 3 Klebsiellae; Klebsiellae *pneumonia*, Klebsiellae *oxytoca* and Klebsiellae *ozaenae* as well as 7 srains of *E. coli* including *E. coli* O 55, *E. coli* O 111,

E. coli O ₁₁₅, *E. coli* O ₁₁₉, *E. coli* O ₁₂₅, *E. coli* O ₁₄₆ and *E. coli* O ₁₆₆ had been analyzed and Sodium Deodecyl Sulphate Polyacrylamide Gel Electrophoresis (SDS-PAGE) profiles of these strains were determined. This powerful technique allows very high resolution of protein and has permitted the identification of multiple major proteins components of different molecular weights.

SDS-PAGE analysis of Klebsiellae and *E. coli* showed that they contained from 5 to 12 bands after staining with Commiassie brilliant blue method which molecular weight ranged from 7.218 to 170.97 KDa.

It was obvious that the molecular weight of *E. coli* O_{55} ranged from 134.55 to 7.2118 KDa. The molecular weight of *E. coli* $_{111}$ ranged from 134.55 to 32.208 KDa. The molecular weight of *E. coli* O_{115} ranged from 134.55 to 33.311 KDa. The molecular weight of *E. coli* O_{119} ranged from 131.24 to 8.4471 KDa. The molecular weight of *E. coli* O_{125} ranged from 118 to 28.562 KDa. The molecular weight of *E. coli* O_{146} ranged from 151.11 to 11.741 KDa. The molecular weight of *E. coli* O_{146} ranged from 154.41 to 21.137.

On the same hand the molecular weight of *Klebsiella oxytoca* ranged from 157.72 to 15.859 KDa. The molecular weight of *Klebsiella pneumoniae* ranged from 167.66 to 22.15 KDa. The molecular weight of *Klebsiella ozaenae* ranged from 170.97 to 38.016 KDa. It was noticed that there was a shared band in molecular weight of (E. coli O₁₁₁, lane 2), (E. coli O₁₁₅, lane 3), (E. coli O₁₄₆, lane 6), (E. coli O₁₆₆, lane 7) and (*Klebsiella ozaenae*, lane 10) in 102.56 KDa , (E. coli O₁₂₅, *lane 5) and*(*Klebsiella oxytaca, lane 8*) *in 103.56KDa and* (*E.* coli O₅₅, lane 1), (E. coli O₁₁₁, lane 2) and (*E. coli* O₁₁₅, lane 3) in 74.588. On the other Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

hand, there was sharing to large extant in molecular weight of band between (E. coli O₁₁₁, lane 2) 42.647KDa, (E. coli O₁₁₉, lane 4) 42.937 KDa and (E. coli O₁₂₅, lane 5) 42.647 KDa and between (E. coli O₁₄₆, lane 6) 40.911 KDa and (E. coli O_{166} , lane 7) 40.621 KDa and finally between (*Klebsiella oxytoca*, lane 8) 41.489 KDa and (*Klebsiella* ozaenae, lane 10) 41.489 KDa. Also there was sharing to large extant in molecular weight of other band which was observed in (E. coli O₅₅, lane 1) 32.576 KDa and (*E. coli* O_{111} , lane 2) 32.208 KDa. and between (*E. coli* O₁₁₅, lane 3) 33.311 KDa. and (*E. coli* O₁₁₉, lane 4) 33.678 KDa. and between (E. coli O₁₄₆, lane 6) 34.956 KDa. and (E. coli O₁₆₆, lane 7)) 34.414 KDa. and finally between (E. coli O₁₂₅, lane 5) 36.568 KDa. and (Klebsiella ozaenae, lane 10) 36.016. It was noticed that (E. coli O₅₅, lane 1), (*E. coli* O₁₁₁, lane 2), (*E. coli* O₁₁₅, lane 3) and (*E. coli* O₁₁₉, lane 4) shared in major crude protein 134.55, 134.55 and 134.55 KDa. This greed to some extent with *Durrani et al.* (2008) who recorded that the major molecular weights of *E. coli* strains were (97.4 and 133 KDa). and minor molecular weight (18, 25 and 48 KDa). It could be primarly differentiate between examined E. coli and Klebsiellae with SDS. No clear pattern could be established between specific serotypes of *E. coli* in relation to SDS bands.

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Kafrelsheikh Vet. Med. J. Vol. 9 No. 1 (2011)

الملخص العربى

ان الجمال تستخدم فى انتاج اللحوم والألبان والنقل الى مناطق صعب الوصول اليها والالتهاب الرئوى فى الجمال مشكلة تؤثر على انتاج الحيوان وتسبب خسائر خطيرة على اقتصاد الحيوان.

اجريت هذه الدراسة على 308 عينة من رئات الجمال المذبوحة منهم 205 عينة رئات سليمة ظاهريا و 103 عينة مصابة وقد وجد ان203 عينة بنسبة 65.91% ايجابية للغزل البكتيرى منهم 124 عينة بنسبة بنسبة فاهريا وكانت 79 عينة بنسبة عينة بنسبة عينة بنسبة 104% وكانت 79 عينة بنسبة عينة بنسبة عينة بنسبة مصابة وقد وجد ان البكتيرى من رئات سليمة ظاهريا وكانت 79 عينة بنسبة عينة بنسبة عينة بنسبة 104% وكانت 79 عينة بنسبة عينة بنسبة مصابة وقد وجد ان البكتيرى من رئات سليمة ظاهريا وكانت 79 عينة بنسبة عينة بنسبة عينة بنسبة 100% ايجابية للعزل البكتيرى من رئات سليمة ظاهريا وكانت 79 عينة بنسبة 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 104 بنسبة 2015% بينما تتائية العزل كانت 79 هذه بنسبة 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 200 بنسبة 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 200% من رئات 200% من رئات مصابة وقد وجد ان العترات البكتيرية فردية العزل 200% من رئات 200% من رئات 200% من ما رئات المليمة التى كانت 50 بنسبة 200% من رئات الميمة التى كانت 50 بنسبة 200% من رئات مليمة ظاهريا التى كانت اعلى من ثتائية العزل عن الرئات السليمة التى كانت 50 بنسبة 200% اعلى فى المصابة عن مليما العكس فى هذه العينات كانت ثتائية العزل وهى 40 بنسبة 200% اعلى فى المصابة عن فردية العزل فى الرئات المصابة وهى 20%.

وقد وجد ان العدد الكلى للمعزولات البكتيرية 302 منهم 174 من رئات سليمة ظاهريا و 128 من المعزولات من رئات مصابة وأن العدد الكلى للبكتيريا السالبة لصبغة الجرام 170 بنسبة 56.29% منهم94 بنسبة 54.02% و 76 بنسبة 59.38% من رئات سليمة ظاهريا ومصابة على التوالى كما يتضح فى جدول (3).

تبين بوضوح ان اكثر المعزولات الفردية العزل وسالبة لصبغة الجرام هى معزولات الميكروب العصوى القولونى من 28 عينة بنسبة 26.92% والسودمونس ايروجونزا 9 عينات بنسبة 8.65 والستروباكتر فيروندى 7 عينات بنسبة 6.73% والكلبسيلا نيمونى 6 عينات بنسبة 5.77 والستروباكتر 5 عينات (4.81%) والكلبسيلا اوكستوكا 3عينات بنسبة 2.89% ومانهيمانيا هيمولتيكا ويرسينيا انيروكولتيكا 2 عينة (1.92%) والكلبسيلا اوزونى والباستريلا ملتوسيدا عينة واحدة (0.96%).

اما بالنسبة للميكروبات ثنائية العزل من رئات الجمال المذبوحة كانت اكثرها حدوثا بروتيس بنييري مع الميكروكوكس بنسبة 13.13% يليه الميكروب العنقودى الذهبى مع القولونى الذى تساوى فى نسبتة مع الميكروب العنقودى الذهبى مع الانتروباكتر والعنقودى الذهبى مع الميكروب السبحى 10.10%.

وقد تم التصنيف السيرولوجى من رئات السليمة ظاهريا والمصابة من الجمال المذبوحة وذلك لمكروب القولونى يشمل على 29 من العزلات البكتيرية من رئات سليمة ظاهريا وهى كالتالى 20 (E. coli O166, E. coli O146, E. coli O119, E. coli O115, E. coli O111) بينما كانت 20 من العزلات البكتيرية من رئات مصابة وكانت كالتالى :

(E. coli O₅₅, E. coli O₁₁₁, E. coli O₁₁₅, E.coli O₁₁₉, E. coli O₁₂₅, E. coli O₁₄₆, E. coli O₁₆₆₎

ولقد وجد من اختبار الضراوة لبعض العزلات البكتيرية التى تم حقنها فى الفئران موت الفئران بنسبة من 20-40-60-100% حيث كانت نسبة الضراوة 100% لميكروب دبلوكوكس نيمونىواركانوبكتيريم بيوجنس والكلبسيلا نيمونى والميكروب المتكور السبحى والسودمونس ايروجونزوا والباستريلا ملتوسيدا ومانهيمانيا هيمولتيكا بينما كانت اقلهم ضراوة هما الميكروب المتكور السبحي والبروتيس فالجاريس وكانت نسبتهم 20% بينما ميكروب الستروباكتر والبروتيس بنيرى لم تسبب اى وفاة فى الفئران.

ولقد تم عمل اختبار الحساسية للمضادات الحيوية بالنسبة للميكروبات المعزولة وتبين ان كلوكسيسلين ودوكسيسيكلين والانروفلوكساسين والجنتاميسين والنوروفلوكساسين هى الاكثر تاثيرا على ميكروب اركانوبكتيريم بيوجينس بينما السيبروفلوكساسين وكلوكساسيلين والانروفلوكساسين والجنتاميسين والنوروفلوكساسين كانت اكثر المضادات الحيوية تاثيرا على دبلوكوكس نيمونى والميكروكوكس والميكروب السبحى والعنقودى الذهبى.

ووجد ان الكلورامفينكول والسيبروفلوكساسين والانروفلوكساسين وفلوموكوين والجنتاميسين والنوروفلوكساسين اكثر المضادات الحيوية التى تؤثر على الستروباكتر فيروندى والميكروب العصوى القولونى والانتروباكتر والكلبسيلا اكستوكا والكلبسيلا اوزونى والكلبسيلانيمونى

السيبروفلوكساسين ودوكسى سيكلين والانروفلوكساسين والنروفلوكساسين اكثر المضادات الحيوية التى تؤثر على مانهيمانيا هيمولتيكا والباستريلا ملتوسيدا ووالبروتيس بنبيري والبروتس فالجلريس والسودمونس ايروجونزا ويرسينيا انتروكولتيكا.

تم اجراء التحليل الكهروكيمائيى باستخدام اس دى اس ل 10 عينات من ميكروب الكلبسيلا والميكروب القولونى واتضح انها تحتوى من 5-12 رابطة بعد صبغها بصبغة الكماسى الزرقاء والتى كان الوزن الجزيئى يتراوح من 7.218 الى 170.97 كيلو دالتون.

وقد وجد ان الوزن الجزيئى لميكروب القولونى E. coli O₅₅ يتراوح من 7.2118 الى 7.2115 الى 134.55 كيلو دالتون كيلو دالتون اما بالنسبة لميكروب O₁₁₁ ولام د coli O₁₁₁ يتراوح من 32.208 الى 134.55 كيلو دالتون وبالنسبة لميكروب I34.55 ولامى تتراوح من 33.311 الى 134.55 كيلو دالتون ولميكروب E. coli O₁₁₅ كيلو دالتون وبالنسبة لميكروب coli O₁₁₅ ولامى تتراوح من 23.311 الى 134.55 كيلو دالتون ولميكروب . وبالنسبة لميكروب S. coli O₁₂₅ كيلو دالتون وبالنسبة لميكروب I34.55 كيلو دالتون الميكروب من 28.562 الى 118 كيلو دالتون وبالنسبة لميكروب E. coli O₁₄₆ الى 11.741 الى من 28.562 الى 118 كيلو دالتون وبالنسبة لميكروب E. coli O₁₄₆ من 11.741 الى

وعلى الجانب الاخر كان الوزن الجزيئى اكلبيسيلا اوكستوكا يتراوح من 15.859 الى 157.72 اما بالنسبة لكلبسيلا نيمونى فالوزن الجزيئى يتراوح من 22.15 الى 167.66 واخيرا الوزن الجزيئى لكلبسيلا اوزونى فهى تتراوح من 38.016 الى 170.97 كيلو دالتون.