

Some bacterial caused abortion in bovine in some governorates of Egypt.

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Abstract

A total of 100 samples of aborted animals 60 buffaloes and 40 cows which proved to be negative for brucellosis by Rose Bengal test were examined bacteriologically for some bacteria other than brucella during the period from May 2011 to June 2012. Samples were collected from private farms in El- Menoufeya, Gharbia, Kalubia and Giza governorates.

The bacteriological examination of aborted foeti, vaginal discharges and placenta of aborted buffaloes revealed the isolation of *Arcanobacterium pyogenes*, *Escherichia coli*, *Staphylococcus aureus*, *Salmonella enteric serovar Dublin*, *Salmonella enteric serovar Typhimurum*, *Campylobacter fetus* subsp. *fetus*, *Campylobacter. fetus* subsp. *Venerealis*, *Streptococcus spp.* and *Pseudomonas aeruginosa* with incidences of 33.3%, 20%, 16.67%, 11.67%, 8.3%, 11.67%, 5%, 6.67% and 3.33%, respectively. While the bacteria isolated from aborted cows were *Arcanobacterium pyogenes*, *Escherichia coli*, *Staphylococcus aureus*, *Salmonella enteric serovar Dublin*, *Salmonella enteric serovar Typhimurum*, *Campylobacter. fetus* subsp. *fetus*, *Campylobacter. fetus* subsp. *Venerealis*, *Streptococcus spp.* and *Pseudomonas aeruginosa* with incidences of 40%, 22.5%, 17.5%, 12.5%, 10%, 12.5%, 7.5%, 7.5% and 5%, respectively.

In vitro antibiogram test indicated that, some tested strains of *Arcanobacterium pyogenes* were highly sensitive to amoxycillin, ciprofloxacin and enrofloxacin. As regards to *E. coli* and *Salmonella* spp isolates, the majority of strains were highly sensitive to cephalocin, erythromycin and gentamicin, Strains of *S. aureus* were highly sensitive to cephalothin, and erythromycin. Meanwhile *Campylobacter fetus* subsp. *fetus* and *Campylobacter fetus* subsp. *Venerealis* were highly sensitive to enrofloxacin, erythromycin, norfloxacin, ciprofloxacin and gentamicin.

Using PCR technique to confirm the results, in case of *Arcanobacterium pyogenes* the molecular bands appeared at 270 bp, while in *SopE* gene of *Salmonella enteric serovar Dublin*, the molecular bands appeared at 399 bp.

Key words: *salmonella*, Abortion, *Arcanobacterium pyogen.*

Introduction

Abortion is one of great concern to farmers due to loss of newborn, loss of fertility or even sterility resulting from prolonged uterine diseases and secondary bacterial infection of the genital tract. Also abortions have a highly negative impact on reproductive efficiency, resulting in significant economic losses for the cattle industry (De Vries, 2006).

Bovine abortion may be due to infectious, toxic, endocrine, physical or nutritional causes. Infectious agents associated with abortion in cattle includes viruses, bacteria, protozoa, and fungus. The exact proportion of cases due to infectious agents is not known, but in

90% of cases in which an etiologic diagnosis is achieved the cause is infectious (Nascimento and Santos, 2003 and Anderson, 2007).

Abortions are caused by many bacterial pathogens that are part of the normal microflora or present in the environment (Butachaiah and Khera 1982, Barr and Anderson 1993 and Yaeger Holler 2007). These bacteria might include, *Arcanobacterium pyogenes*, *Bacillus spp.*, *Escherichia coli*, *Pasteurella spp.*, *Pseudomonas spp.*, *Salmonella spp.*, *Staphylococcus spp.*, and *Streptococcus spp.* Bacteria are able to spread hematogenously to the fetoplacental unit and cause abortion, usually late in gestation. *Arcanobacterium pyogenes* capable

of producing suppurative lesions in any organ or tissue in animals. In farm animals, especially ruminants, it is the most common bacteria found in infected wounds and abscesses which is also an important cause of mastitis in cattle. Previously called *Actinomyces pyogenes* and *Arcanobacterium pyogenes* (**Hasan *et al.*, 2005** and **Carlos *et al.*, 2009**)

Early studies on *Arcanobacterium pyogenes* as a cause of bovine abortion was reported by (**Boyd and Kelly 1943**). On farms with records of *Arcanobacterium pyogenes* abortions all endometritis and mastitis cases could be associated with the same infection. The organisms were demonstrated in the uterine secretion in both clinically normal cows and bull semen. This organism was found to be viable for a long time, and, therefore, represented a serious source of infection--abortions, endometritis, and mastitis (**Büchvarova, 1982**). In cattle, it is a common cause of abscesses and mastitis and is often involved with pneumonia (**Blood and Radostits 1990**). The isolation of *Arcanobacterium pyogenes* from the stomach content or fetal tissues was considered diagnostic, because *Arcanobacterium pyogenes* is a common postparturient invader of the uterus, especially when the placenta is retained. Isolation of *Arcanobacterium pyogenes* from only a retained placenta was considered conclusive evidence that *Arcanobacterium pyogenes* caused the abortion only if it was accompanied by suppurative placentitis (**Blood and Radostits 1990**, **Fahriye *et al.*, 2011** and **Manigeh *et al.*, 2011**).

Many abortions may be seen in herds where an outbreak of enteric disease has occurred. Carrier animals and faecal-oral transmission are common. Diagnosis is mainly based on isolation and identification of the agent from placenta and foeti. *Salmonella spp* cause storms abortion, the cows are usually sick and the foeti and placentas are autolyzed and emphysematous. Salmonellae can be isolated from the abomasal contents, foeti tissues, from uterine fluids and the dams' faeces (**The Merck Veterinary Manual, 2011**). *Salmonella enteric serovar Dublin*, is usually associated with en-

teric infection and diarrhea, particularly in calves, although infection of pregnant cows may result in abortion as the only clinical manifestation of infection in the herd (**Ring, 1985**). In cattle, *Salmonella enteric serovar Dublin*, causes economic losses and death among calves and young animals, as well as abortion, reproductive disorders among adult cattle, extra labour and increased veterinary expenses (**Nascimento and Santos 2003** and **Nielsen *et al.*, 2004**). Salmonellosis due to *Salmonella enteric serovar Dublin*, infection in cattle results in high morbidity and mortality in calves and significant morbidity in adult cattle (**Rice *et al.*, 1997**).

E. coli causes bovine abortion so often in aborted foeti that have compatible lesions in the abortions and cannot be ignored. The fact that it is the third most common bacterium implicated in bovine abortions probably can be attributed to its prevalence in the environment of cattle. The aborted foeti were usually moderately to severely autolyzed, and often there was bacterial decomposition with gas bubbles in the tissues and an unpleasant odor. Fibrinous peritonitis and/or pleuritis were often present, as well purulent placentitis and/or pneumonia.. It has been isolated in the case of a partial abortion (**Linde 1983**).

Staphylococcus aureus is a normal inhabitant of the skin and mucus membrane including genital tract (**Corbellini *et al.*, 2006**). Also *Staphylococcus aureus* is commonly isolated from bovine mastitis and occasionally recovered from the uterus. These bacteria can infect the fetus causing sporadic abortion in cattle (**Anderson *et al.*, 1990** and **Kirkbride, 1992**).

PCR has been increasingly used as a diagnostic tool for etiologic diagnosis of abortion in cattle either as a complement or replacement of time consuming traditional diagnostic methods such as bacteria (**Anderson, 2007**). Considering the potential of PCR for etiological diagnosis of infectious bovine abortion, the objective of this study was to use several previously established PCR protocols as a comprehensive PCR panel for identification of etiologic agent in tissues from aborted bovine foeti. This study

included frozen tissues as well as formalin-fixed and paraffin-embedded tissues from aborted foeti and placentas.

PCR protocols have been recently developed for identification of infectious agents in aborted bovine foeti, including *Salmonella enterica* serovas. *Dublin* or *Salmonella* species. (Kerouanton *et al.*, 1996, Amavisit, 2001 and Whyte, 2002), and *Arcanobacterium pyogenes* (Jost,2002), with the exception of *Arcanobacterium pyogenes* and *Salmonella enteric* ser. *dublin*, these agents are primarily involved in reproductive diseases in cattle.

The aim of this work is the evaluation of some bacteriological aetiology caused bovine abortion in some governorates of Egypt. On the other hand, susceptibility of most isolated bacteria to chemotherapeutic agents was done, to overcome the problem and reduce losses.

Material and Methods:

Collection of samples:

The samples were collected under complete aseptic conditions from 100 aborted animals ; (60) buffaloes and (40) cows for bacteriological examination during period from May 2011 till June 2012. Swabs from all examined samples of stomach contents of aborted foeti as well as liver, spleen, and lungs were collected in separated sterile containers and were transported as quickly as possible to the lab in ice box .

Placenta and vaginal discharges were collected aseptically from aborted buffaloes and cows and transferred immediately to the laboratory, where they were examined bacteriologically. Three samples collected from each animal (one of each of aborted foeti, vaginal discharge and placenta) .

All aborted buffaloes and cows were serologically negative to brucellosis when screened by Rose Bengal Plate Test. All samples were obtained from aborted buffaloes and cows from various private farms at El- Menoufeya, Gharbia, Kalubia and Giza governorates.

Bacteriological examination:

The collected samples, (swabs from stomach

(abomasal), intestinal contents, vaginal samples and placenta of the aborted dam as well as internal organs) were inoculated directly in nutrient broth and incubated at 37°C for 6 hours. Loopful from each broth culture were inoculated directly onto nutrient agar, blood agar, Macconkey's agar, bile salt lactose agar, *Salmonella* . Shigilla agar, Mannitol Salt agar, Baird parker's agar media for isolation of *S. aureus*, Eosin Methylene blue agar media for isolation of *E. coli* and *Enterobacteriaceae* family, and Campylobacter blood free selective agar. The plates were incubated at 37°C for 24-48 hours. Suspected colonies onto the surface of these media were identified by studying characters of the colonies as well as Gram's stain, then identified morphologically according to the method described by (Merchant and Packer,1969, Kloss and Schleifer 1986, and Barrow and Feltham 1993).

One single colony showed typical colonial appearance and morphological characters was picked up and streaked into semisolid agar media and incubated at 37°C for 24 hours to obtain pure culture, for further identification. The pure colonies were biochemically identified according to (Cruickshank *et al.*,1975, Koneman *et al.*, 1992 and Quinn *et al.*,2002). The Gram negative bacteria included *Enterobacteriaceae* family (Krieg and Holt, 1984).

The inoculated plates for Campylobacter blood free selective agar supplemented with antibiotics, were incubated in microaerophilic condition using gas generating kit in plastic anaerobic jar at 37 °C for 3- 4 days. Then suspected *Campylobacter* colonies were identified according to Skirrow and Benjamin, (1980) and Prescott and Munroe, (1982).

Susceptibility of isolates to various chemotherapeutic agents :

Antibiotic sensitivity test was applied on pure subcultures according to Finegold and Martin (1982) and Ababneh and Degefa (2006) to detect the drug of choice against different iso-

lated bacteria for trials of treatment of the infected animals. The results were interpreted according to **Koneman *et al.*, (1992).**

Molecular technique:

1- Extraction of DNA from frozen tissue samples:

Extraction of DNA was performed using DNA purification Kit, Promega, USA according to the manufacturer's instructions. Briefly, 100µL of thawed homogenates of fetal tissues were mixed with 600µL of Nuclei Lysis Solution and homogenized for 10 seconds. Samples were incubated at 65°C for 30min, followed by addition of 20 µL proteinase K and incubation at 56°C for 3 hours, vortexing every hour. Three microliters of RNase A (4mg mL⁻¹) were added, the samples were mixed and incubated at 37°C for 30min. After cooling, 200µL of Protein Precipitation Solution were added, followed by vortexing and centrifugation at 13,000 x g for 4min. The supernatant was transferred to a new micro tube with 600µL of isopropanol, mixed, and centrifuged at 13,000x g for 3min. The supernatant was discarded and the pellet was washed with 600µL of 70% ethanol, followed by a final centrifugation at 13,000x g for 3min. Each pellet was dissolved in 100µL of DNA Rehydration Solution by incubating at 65°C for 1 hour. **(Shi., 2004).** 500µL of 0.1M NaOH was added to each micro tube containing two 10µm-thick tissue sections, and heated at 100°C for 20min. Then, 500µL of phenol: chloroform: isopropanol alcohol (25:24:1) were added, followed by vortexing and centrifugation at 10,000 x g for 10 min. The supernatant was transferred to another micro tube and 1 volume of chloroform was added, mixed by vortexing and centrifuged at 10,000x g for 5min. The upper phase of the supernatant was transferred to another micro tube and mixed with 0.1 volumes of 3.0M sodium acetate, and 1 volume of isopropanol, followed by incubation at -20°C overnight. The suspension of precipitated DNA was then centrifuged at 10,000x g for 5 min at 4°C. The pel-

let was washed with 75% ethanol, dried, and diluted in 50µL of distilled water. DNA samples from different tissues of the same fetus were pooled prior to PCR amplification. Samples that did not yield actin amplicon or had DNA concentration lower than 100 ng µL⁻¹ as assessed by spectrophotometry were excluded from further analysis. DNA samples were PCR tested for detection of bacteria infectious agents known to cause abortion in cattle, including *Arcanobacterium pyogenes*, and *Salmonella* sp.

PCR for *Arcanobacterium pyogenes*:

The primers including 5'-GGCCCGAATGT-CACCGC-3' (forward primer) and 5'-AC-TCCGCCTCTAGCGC-3' (reverse primer) were used for PCR amplification of *Arcanobacterium pyogenes*, designed by **(Jost., 2002)**

The PCR mixture (50 µL) contained 5 µL 10x PCR buffer, 6 µL forward primer, 5 µL reverse primer, 1 µL dNTP, 1.5 µL MgCl₂, 0.5 µL Taq polymerase and 2 µL DNA template. The PCR-amplification was performed with temperature program consisting of initial denaturation (94°C, 3 min) and 30 cycles of denaturation (94°C, 1 min), primer annealing (55°C, 1 min) and primer extension (72°C, 2 min). For final extension, this was followed by incubation at 72°C for 10 min. A volume of 5 µL of the PCR product was analyzed by electrophoresis in 1% agarose gel.

PCR for *Salmonella entrica* serovas

Dublin:

The primers including 5'-ACACACTTTCA-CCGAGGAAGCG-3' (forward primer) and 5'-GGATGCCTTCTGATGTTGACTGG-3' (reverse primer) were used for PCR amplification of *SopE* gene, designed by **(Rahman *et al.* 2004).** The PCR mixture (50 µL) contained 5 µL 10x PCR buffer, 6 µL forward primer, 5 µL reverse primer, 1 µL dNTP, 1.5 µL MgCl₂, 0.5 µL Taq polymerase and 2 µL DNA template. The PCR-amplification was performed with temperature program consisting of initial denaturation (94°C, 3 min) and 30 cycles of denatu-

ration (94°C, 1 min), primer annealing (56°C, 1 min) and primer extension (72°C, 2 min). For final extension, this was followed by incubation at 72°C for 10 min. A volume of 5 µL of the PCR product was analyzed by electrophoresis in 1% agarose gel.

RESULTS

Results in Table (1), shows the total bacterial isolates from 60 (aborted foeti, 60 vaginal discharge and 60 placenta) of aborted buffaloes which were (63.3%, 28.3% and 25% respectively). Out of total 180 samples, 70 samples (38.9%) were positive for bacteriological examination.

The bacteria isolated from aborted foeti of buffaloes were *Arcanobacterium pyogenes*, *Escherichia coli*, *Staphylococcus aureus*, *Salmonella enterica serovas Dublin*, *Salmonella enterica serovas Typhimurum*, *Campylobacter. fetus subsp. Fetus*, *Campylobacter. fetus subsp. Venerealis*, *Streptococcus spp.* and *Pseudomonas aeruginosa* with incidence of (16.67%, 11.67%, 8.3%, 6.67%, 5%, 6.67%, 3.33%, 3.33% and 1.67%, respectively). While the bacteriological isolation from vaginal dis-

charge were *Arcanobacterium pyogenes*, *Escherichia coli*, *Staphylococcus aureus*, *Salmonella enteric serovar Dublin*, *Salmonella enteric serovar Typhimurum*, *Campylobacter. fetus subsp. Fetus*, *Campylobacter. fetus subsp. Venerealis*, *Streptococcus spp.* and *Pseudomonas aeruginosa* with incidence of 10%, 5%, 3.33%, 1.67%, 1.67%, 1.67%, 1.67%, 1.67% and 1.67%, respectively. The bacteriological isolation from placenta were *Arcanobacterium pyogenes*, *Escherichia coli*, *Staphylococcus aureus*, *Salmonella enteric serovar Dublin*, *Salmonella enteric serovar Typhimurum*, *Campylobacter. fetus subsp. Fetus*, *Campylobacter. fetus subsp. Venerealis*, *Streptococcus spp.* and *Pseudomonas aeruginosa* with incidence of 6.67% 3.33%, 5%, 3.33%, 1.67%, 3.33%, 0%, 1.67%, and 0%, respectively.

In attempt to correlate the relation between various types of bacteria and their sites of positive aborted buffaloes foeti, the data obtained were recorded in Table (2). Out of 60 aborted buffaloes infected with *Arcanobacterium pyogenes*, *Escherichia coli*, *Staphylococcus aureus*, *Salmonella enteric serovar Dublin*, *Salmonella enteric serovar Typhimurum*, *Campylobacter.*

Table (1): Prevalence of bacteria isolated from vaginal discharge, placenta and aborted foeti of aborted buffaloes.

Microorganisms	Sites of isolation						Total isolates	
	Aborted foeti (60 samples)		Vaginal dis- charge (60samples)		Placenta (60 samples)			
Bacterial isolates	No.	%	No.	%	No	%	No.	%
<i>Arcanobacterium pyogenes</i>	10	16.67	6	10	4	6.67	20	33.3
<i>Escherichia coli</i>	7	11.67	3	5	2	3.33	12	20
<i>Staphylococcus aureus</i>	5	8.3	2	3.33	3	5	10	16.67
<i>Salmonella entrica serovas Dublin</i>	4	6.67	1	1.67	2	3.33	7	11.67
<i>Salmonella entrica serovas Typhimurum</i>	3	5	1	1.67	1	1.67	5	8.3
<i>Campylobacter. fetus subsp. Fetus</i>	4	6.67	1	1.67	2	3.33	7	11.67
<i>Campylobacter. fetus subsp. Venerealis</i>	2	3.33	1	1.67	0	0	3	5
<i>Streptococcus spp.</i>	2	3.33	1	1.67	1	1.67	4	6.67
<i>Pseudomonas aeruginosa</i>	1	1.67	1	1.67	0	0	2	3.33
<i>Total bacterial isolates</i>	38	63.3	17	28.3	15	25	70	38.9

Table (2): Bacteria isolated from aborted buffaloes foeti as regarded to its sites of isolation from internal organs.

<i>Microorganisms</i>	Total isolates	<i>Sites of isolation</i>							
		Stomach content		Liver		Spleen		Lungs	
Bacterial isolates	No.	No.	%	No.	%	No.	%	No.	%
<i>Arcanobacterium pyogenes</i>	10	5	50	3	30	1	10	1	10
<i>Escherichia coli</i>	7	4	57.1	2	28.6	1	14.3	0	0
<i>Staphylococcus aureus</i>	5	2	40	1	20	1	20	1	20
<i>Salmonella</i> entrica serovas Dublin	4	3	75	1	25	0	0	0	0
<i>Salmonella</i> entrica serovas Typhimurum	3	2	66.7	1	33.3	0	0	0	0
<i>Campylobacter. fetus</i> subsp. <i>fetus</i>	4	2	50	1	25	1	25	0	0
<i>Campylobacter. fetus</i> subsp. <i>Venerealis</i>	2	1	50	1	50	0	0	0	0
<i>Streptococcus spp.</i>	2	1	50	1	50	0	0	0	0
<i>Pseudomonas aeruginosa</i>	1	0	0	0	0	0	0	1	100
Total bacterial isolates	38	20	52.6	11	28.9	4	10.5	3	7.9

Table (3): Prevalence of bacteria isolated from vaginal discharge, placenta and aborted foeti of aborted cows.

Microorganisms	Sites of isolation						Total isolates	
	Aborted foeti (40)		Vaginal discharge (40)		Placenta (40)			
A-Bacterial isolates	No.	%	No.	%	No.	%	No.	%
Arcanobacterium pyogenes	8	20	5	12.5	3	7.5	16	40
Escherichia coli	5	12.5	2	5	2	5	9	22.5
Staphylococcus aureus	4	10	1	2.5	2	5	7	17.5
Salmonella Dublin	3	7.5	1	2.5	1	2.5	5	12.5
Salmonella Typhimurum	2	5	1	2.5	1	2.5	4	10
Campylobacter. fetus subsp. Fetus	3	7.5	1	2.5	1	2.5	5	12.5
Campylobacter. fetus subsp. Venerealis	1	2.5	1	2.5	1	2.5	3	7.5
Streptococcus spp.	0	0	2	5	1	2.5	3	7.5
Pseudomonas aeruginosa	0	0	1	2.5	1	2.5	2	5
Total bacterial isolates	26	65	15	37.5	13	32.5	54	45

Table (4): Bacteria isolated from aborted cows foeti as regarded to its sites of isolation from internal organs.

<i>Microorganisms</i>	Total isolates	<i>Sites of isolation</i>							
		Stomach content		Liver		Spleen		Lungs	
Bacterial isolates	No.	No.	%	No.	%	No.	%	No.	%
<i>Arcanobacterium pyogenes</i>	8	6	75	1	12.5	1	12.5	0	0
<i>Escherichia coli</i>	5	2	40	1	20	1	20	1	20
<i>Staphylococcus aureus</i>	4	1	25	1	25	1	25	1	25
<i>Salmonella</i> entrica serovas <i>Dublin</i>	3	2	66.7	1	33.3	0	0	0	0
<i>Salmonella</i> entrica serovas <i>Typhimurum</i>	2	1	50	1	50	0	0	0	0
<i>Campylobacter. fetus</i> subsp. <i>Fetus</i>	3	2	66.7	1	33.3	0	0	0	0
<i>Campylobacter. fetus</i> subsp. <i>Venerealis</i>	1	1	100	0	0	0	0	0	0
Total bacterial isolates	26	15	57.7	6	23.1	3	11.5	2	7.8

Table (5): Antibiotic sensitivity test of the different bacterial isolated strains from aborted buffaloes and cows.

Antibacterial agents	Concentration	<i>Arcanobacterium Pyogenes</i> (36)*		<i>E.coli</i> (21)		<i>S. aureus</i> (17)*		<i>Salmonella</i> spp. (21)*		<i>C.fetus</i> supsp. <i>Fetus</i> (12)*		<i>C.fetus</i> supsp. <i>Venerealis</i> (6)*	
		S.	%	S.	%	S.	%	S.	%	S.	%	S.	%
Ampicillin	10ug	3/36	8.3	16/21	76.2	12/17	70.5	17/21	80.9	3/12	25	3/6	50
Amoxycillin	10ug	36/36	100	8/21	38.1	9/17	52.9	7/21	33.3	8/12	66.7	5/6	83.3
Cephalocin	10ug	13/36	41.7	21/21	100	16/17	94.1	20/21	95.2	10/12	83.3	4/6	66.7
Ciprofloxacin	30ug	30/36	83.3	4/21	19	11/17	64.7	8/21	38.1	11/12	91.7	5/6	83.3
Enrofloxacin	30ug	30/36	83.3	0	0	14/17	82.4	6/21	28.6	12/12	100	6/6	100
Erythromycin	10ug	18/36	50	21/21	100	16/17	94.1	18/21	85.7	11/12	91.7	5/6	83.3
Flumequine	10ug	28/36	77.8	0	0	11/17	64.7	5/21	23.8	5/12	41.7	2/6	33.3
Gentamicin	10ug	3/36	8.3	21/21	100	3/17	17.4	21/21	100	10/12	83.3	5/6	83.3
Norfloxacin	30ug	0/36	0	19/21	90	0/17	0	20/21	95.2	11/12	91.7	5/6	83.3
Polymyxin	30 ug	6/36	16.6	12/21	57.1	0/17	0	14/21	66.7	5/12	41.7	2/6	33.3
Streptomycin	10 U	0/36	0	12/21	57.1	2/17	11.8	15/21	71.4	0/12	0	1/6	16.7
Tetracycline	10ug	0/36	0	8/21	38.1	5/17	29.4	9/21	42.9	2/12	16.7	1/6	16.7
Trimethoprim& Sulphamethoxazole	30ug	26/36	72.2	8/21	38.1	12/17	70.5	10/21	47.6	6/12	50	3/6	50
Tobramycin	1.25 ug	27/36	75	21/21	100	15/17	88.2	19/21	90.5	2/12	16.7	1/6	16.7

* Number of isolates. %: Percentage of sensitive isolates in relation to total isolates.

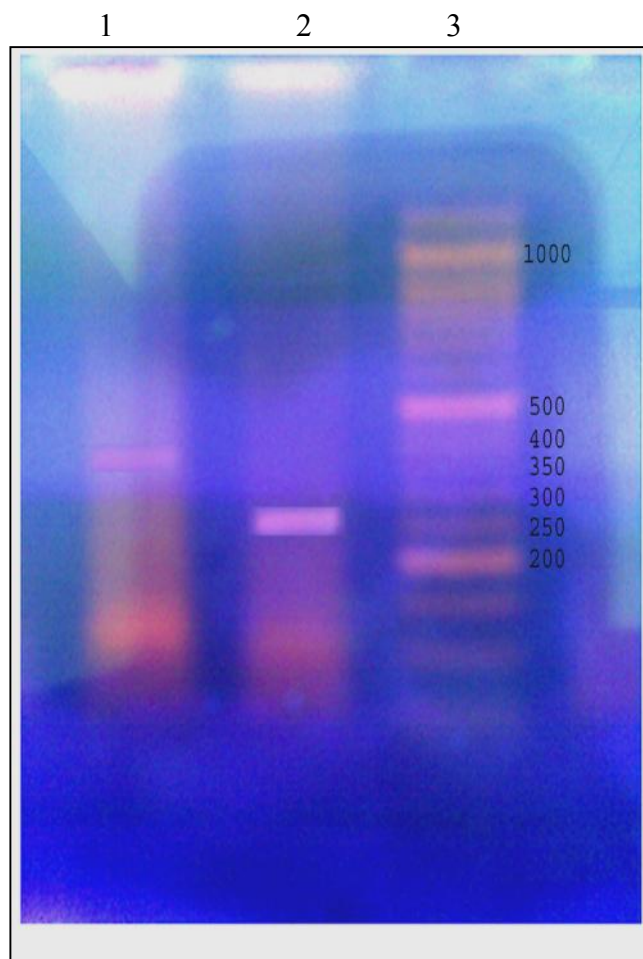


Fig (1) : Compare the *sopE* gene extracted from *S. enterica* serovas *Dubline* and *Arcanobacterium pyogenes* with DNA marker.

lane1: PCR product of the *SopE* gene at 399 bp

Lane 2: *Arcanobacterium pyogenes* at 270 bp

Lane 3: DNA marker (GeneDirex Cat No.DM012-R500)

Discussion

Abortion is one of the most important world-wide factors and its economical losses is the predominant index in dairy cattle breeding.

In this study, the bacteriological examination of 60 of (aborted foeti, vaginal discharge and placenta) of aborted buffaloes revealed that 70 samples (38.9%) were positive for bacteriological examination. While out of 40 of (aborted foeti, vaginal discharge and placenta) of aborted cows revealed that 54 samples (45%) were positive for bacteriological examination. These results are nearly agree with

Hubbert, *et al.*, (1973), Kirkbride, *et al.*, (1973), Miller, (1977) and Miller (1987), as they reported the percentage of bacterial causes of bovine abortion was 42.6%. While **Jamaluddin *et al.* (1996)** recorded that bacterial abortion accounted for 18% of the total etiological diagnosis.

As shown in **Tables (1 & 3)**, the bacteriological examination illustrated that the *Arcanobacterium pyogenes* was the most isolated microorganism from aborted buffaloes and cows with an incidence of 33.3% and 40% respectively. They were mostly isolated from aborted buffaloes foeti, vaginal discharge and placenta

with the incidence of 16.67%, 10% and 6.67% respectively. Meanwhile isolated from aborted cows foeti, vaginal discharge and placenta with the incidence of (20%, 12.5% and 7.5% respectively). Then followed by *Escherichia coli*, *Staphylococcus aureus*, *Salmonella entrica serovas Dublin*, *Salmonella entrica serovas typhimurum*, *Campylobacter. fetus* subsp. *Fetus*, *Campylobacter. fetus* subsp. *venerealis*, *Streptococcus* spp. and *Pseudomonas aeruginosa* with incidence of (20%, 16.67%, 11.67%, 8.3%, 11.67%, 5%, 6.67% and 3.33%, respectively), from aborted buffaloes foeti, vaginal discharge and placenta. But followed by *Escherichia coli*, *Staphylococcus aureus*, *Salmonella entrica serovas Dublin*, *Salmonella entrica serovas Typhimurum*, *Campylobacter. fetus* subsp. *fetus*, *Campylobacter. fetus* subsp. *Venerealis*, *Streptococcus* spp. and *Pseudomonas aeruginosa* with incidence of (22.5%, 17.5%, 12.5%, 10%, 12.5%, 7.5%, 7.5 % and 5%, respectively), from aborted cows foeti, vaginal discharge and placenta. These results agree with **Erin et al. (2005)** who showed that the most common and economically important bacteria for uterine infection were *Actinomyces*, *Escherichia coli*, *Fusobacterium*, *Pasteurella*, *Pseudomonas* and *Staphylococcus*. In other study by **Gani et al. (2008)** showed that *Staphylococcus* spp was predominant (37.8%), followed by *E.coli* (29.7%), *Pseudomonas* (18.9%), while Gram negative minute rod shaped bacteria was found in 24.3% of cases in repeat breeder cows. While **Foldi, et al. (2006)** recorded that, *Arcanobacterium*, *Staphylococcus*, *Streptococcus*, *E. coli* and *Klebsiella* spp are found to be associated with endometritis in cows. Also **Anderson et al. (1990)**, found *Actinomyces pyogenes*, *Streptococcus* spp., *Escherichia coli*, and *Bacillus* spp., which are regarded as causes of sporadic bovine abortion. While **Ebrahim et al. (2012)**, reported that all the tested aborted bovines were bacteriologically positive to *E.coli* (39%), *Staphylococcus aureus* (13%), *Staphylococcus epidermicus* (9.5%), *Streptococcus faecalis* (9.5%), and *Bacillus* (13%).

Specimens required for laboratory diagnosis of

bovine abortions are abomasal contents, liver, spleen and lungs after faetal death. Clarifying the role of bacteria in aborted foeti may lead to more extensive light involving problem of bovine abortion. Distribution of bacteria species in aborted dead foeti and its incidence is represented in Table (2 and 4). It is worthy to mention that stomach of aborted foetus harboured most isolated bacteria from dead aborted foeti and the stomach contents was the most common seat for isolation and considered as the specimen of laboratory choice. It was followed by liver and spleen specimens. While the lungs specimens were the least organs infected by bacteria from examined aborted foeti. These findings coincide with the observations obtained by **Hunter et al., (1976)** who recorded isolation of *Salmonella enteric serovar Typhimurum* from fetal stomach contents and placenta. **Andreson et al., (1983)** isolated of *Campylobacter* spp. from stomach contents of goats. Also this finding was in agreement with several other studies **Jerrett et al., (1984)** and **Kirkbride, (1992)**, who indicated that in situations where specimen submission is optimal, where most submissions consisted of fetus, placenta, and maternal blood or their combinations, a high rate of achieving definitive diagnosis may be obtained. **Jamaluddin et al., (1996)**, submissions comprising fetus, placenta, maternal blood, or their combinations were associated with a higher likelihood of definitive diagnosis for abortion than tissues. Table (5) illustrated in vitro sensitivity of the most prevalent bacteria isolated from aborted buffaloes and cows to (14) chemotherapeutic agents. All tested strains of *Arcanobacterium pyogenes* were highly sensitive to amoxycillin, ciprofloxacin, enrofloxacin, flumequine and tobramycin. These findings are nearly similar with that reported by **Zaki (1999)** and who revealed that the majority of *Arcanobacterium pyogenes* isolated from opened skin lesions in buffaloes were highly sensitive to enrofloxacin, ciprofloxacin, ampicillin and tobramycin. As regards to *E. coli* and *Salmonella* spp isolates, the majority of strains were highly sensitive to cephalocin,

erythromycin, gentamicin, tobramycin and norfloxacin. These agree with the results of (Mishra ., 1992). All tested strains of *Staph. aureus* were highly sensitive to cephalothin, erythromycin and tobramycin. These results are nearly in agreement with that of Havelka (1983), and Hazarika *et al.*, (1991). Meanwhile enrofloxacin, erythromycin, norfloxacin, ciprofloxacin and gentamicin were the most effective anti bacteria for the *Campylobacter fetus* subsp. *featus* and *Campylobacter fetus* subsp. *venerealis*. These findings are in agreement with that obtained by (Zenin 1985 and Narita *et al.*, 1988)..

One of the objectives of this study was to investigate the availability of PCR methods for the detection of bacterial infection. PCR assay is an effective method for the detection of wide range of genes and biological agents. This technique does not have the described limitations in the other methods. Therefore, these described time consuming or expensive procedures for identification of pathogen can be replaced with a PCR reaction (Eisa *et al.*, 2011). The results obtained in this study showed the rapid and successful use of PCR in the identification of doubtful samples in comparison with traditional microbiological culture methods that may take several days to complete.

In our study of *Arcanobacterium pyogenes*, PCR using DNA template of the isolates and specific primers was used for further confirmation of the pathogen. PCR products were analyzed using agarose gel electrophoresis and purified from the agarose gel. *Arcanobacterium pyogenes* is currently widespread pathogens throughout the world and has prompted a heightened interest and concern for the rapid detection of these pathogens. Thus, rapid and accurate detection approaches are needed to reduce risk infection caused by *Arcanobacterium pyogenes*. This study was aimed to establish simple and rapid testing methods based on biochemical test for identification of *Arcanobacterium pyogenes*. No false positive amplification was observed, indicating the high specificity of the established PCR assay.

PCR reaction was performed and specific band of PCR product was observed at the position corresponding to the expected size of the DNA amplification products about 399 bp for *SopE* gene and at 270 bp for *Arcanobacterium pyogenes* compare with DNA ladder GeneDirex. The PCR product was checked by agarose gel electrophoresis and stained with ethidium bromide as shown in (Fig. 1). Due to concentrations of genomic DNA in the sample that was below the limit of detection of the traditional method. Therefore, these results indicate the potential of PCR as an additional diagnostic tool for identification of infectious agents causing bovine abortion. Importantly, when several diagnostic approaches are employed together, the rate of success apparently is increased in comparison to PCR alone (Anderson, 2007). It is noteworthy that extremely high sensitivity of PCR methods under laboratorial conditions may not be reproducible under field conditions since suboptimal sampling and storage may have a major impact on sensitivity as recently reported for the application of a diagnostic PCR-based method under field conditions in Brazil (Paixao, 2008).

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