



A PRELIMINARY STUDY ON THE ISOLATION AND IDENTIFICATION OF AEROBIC BACTERIA FROM DIABETIC FOOT INFECTION AND ITS ANTIBIOGRAM

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ABSTRACT

Microbiology study in diabetes is metabolic disorder affecting the Langer islets of pancreas affecting the insulin and glucagon production and in turn affecting the sugar (glucose) metabolism which in turn affecting the overall metabolism the disease may be classified in type I diabetes mellitus and type II diabetes mellitus. Type I diabetic mellitus It is formerly known as insulin- dependent diabetic mellitus type II diabetes mellitus It is characterized by hyperglycemia due to an individual's resistance to insulin with an insulin secretory defect. Treatment the oral hypoglycemic agents are sulphonyl urea and biguanides. The sulphonyl urea drugs (tolbu – tamide and glibenclamide) stimulate insulin secretion. Diabetic foot infections are infections that can develop in the skin, muscles or bones of the foot as a result of nerve damage and poor circulation that is associated with diabetes. Diabetes mellitus is a disorder that primarily affects the microvascular circulation. Collection of sample diabetic patients included in this study was of age group (35-45). The foot sample of both sexes were collected from government hospitals in Cheyyar, a sterile swab was used for collecting sample from diabetic patients study in sample processing, Microscopic examination, Antibiotic sensitivity test, Statistical analysis, the samples were processed according to the standard microbiological techniques the aerobic organisms were isolate from pus culture statistical analysis was done. *Staphylococcus aureus* was the predominant organism among the isolation the gram negative organism *Pseudomonas aeruginosa* was predominant other organism included in our study was *Escherichia coli* and *Proteus mirabilis*.

Keywords: Diabetic foot infection, Antibiotic sensitivity test, Microscopic examination, Sample processing.

INTRODUCTION

Diabetes is metabolic disorder affecting the Langer islets of pancreas affecting the insulin and glucagon production and in turn affecting the sugar (glucose) metabolism which in turn affecting the overall metabolism the disease may be classified in type I diabetes mellitus and type II diabetes mellitus. Type I diabetic mellitus It is formerly known as insulin- dependent diabetic mellitus type II diabetes mellitus It is characterized by hyperglycemia due to an individual's resistance to insulin with an insulin secretory defect [1]. The resistance results in a relative. The changes are mainly the result of

low insulin glucagons ratio. The effects of insulin on carbohydrate lipid and amino acid metabolism have been described in diabetic Mellitus all these effects are reversed [2].

Treatment the oral hypoglycemic agents are sulphonylurea and biguanides. The sulphonyl urea drugs (tolbu – tamide and glibenclamide) stimulate insulin secretion [3]. They are used in Type -2 disease. Insulin is injected as replacement therapy in Type 1 disease, as well as in Type 2 disease. Where oral drugs are not sufficient. There are many hypoglycemic drugs, which belong to the group sulfonamide, are tolbutamide (Orinase),

chlorpropamide (diabinese) and tolazamide (tolinase) [4].

Diabetic foot infections are infections that can develop in the skin, muscles or bones of the foot as a result of nerve damage and poor circulation that is associated with diabetes. Diabetes mellitus is a disorder that primarily affects the microvascular circulation [5]. In the extremities, microvascular disease due to "sugar-coated capillaries" limits the blood supply to the superficial and deep structures. Pressure due to ill-fitting shoes or trauma further compromises the local blood supply at the microvascular level, predisposing the patient to infection. The infection may involve the skin, soft tissues, bone, or all of these tissues [6].

MATERIALS AND METHODS

Collection of sample diabetic patients included in this study was of age group (35-45). The foot sample of both sexes were collected from government hospitals in Cheyyar, a sterile swab was used for collecting sample from diabetic patients. The sample was collected by simply rolling the tip of the swab on its side for one full rotation over the infected area. Dried surface was premoistened with a saline swab which improves the yield, transportation of sample. The collected swab was placed in Stuart's media and transported to the lab.

Processing Of Sample

Microscopic examination, Staining method, Hanging drop method, Culture, Biochemical parameter, Catalase, Oxidase, Coagulase, Indole, Methyl red, Voges-Proskauer, Citrate, Triple sugar iron agar, Urease, Gelatin Hydrolysis, Nitrate reduction, Sugar fermentation, Selective media, Antibiotic sensitivity study. All the analysis are carried out by the method of Sigma Diagnostic kits (Sigma Chemical Company Catalogue, 1997) [7].

Statistical Analysis

All the data were analyzed as per the method of Pillai and Sinha HC. (1968) [8].

RESULTS AND DISCUSSION

Plate 1, Tables 1-10 and Fig 6 indicate the results obtained in the present investigation. Diabetic patients 25 sample had been collected from Government Hospital in Cheyyar. 16 sample were collected from male and 9 sample were collected from female. The number of different isolates from total number of specimens and microscopic examination such as staining motility test and cultural characters, colony morphology, biochemical characters in different isolation specimen in the microorganism for *Staphylococcus aureus*, *Pseudomonas aeruginosa*, *E. coli* and *Proteus mirabilis* identified.

The isolated bacteria were tested for their susceptibility to different antibiotics. *Staphylococcus aureus* were found to be highly [9], sensitive to ampicillin,

amikacin, gentamicin, vancomycin and resistant to methicillin. *Pseudomonas aeruginosa* were found to be highly [10], sensitive to amikacin, gentamicin, ceftazidime and moderately sensitive to ciprofloxacin. *Proteus mirabilis* highly sensitivity to ciprofloxacin and moderately sensitive to tetracycline. It was found to be ampicillin, amoxycillin, cefotaxime, *Escherichia coli* highly tetracycline and moderately [11], sensitive to ampicillin, streptomycin and polymyxin-B. Our study demonstrated the large number and variety of organism can be isolated from properly obtained specimens that are optimally processed.

Samples were collected from the nearby hospital. The samples were processed according to the standard microbiological techniques. The aerobic organisms were isolated from pus culture. Statistical analysis was done. *Staphylococcus aureus* was the predominant organism among the isolation. The gram-negative organism *Pseudomonas aeruginosa* was predominant. Other organism included in our study was *Escherichia coli* and *Proteus mirabilis* [12].

Plate. 1 Diabetic Foot Infection



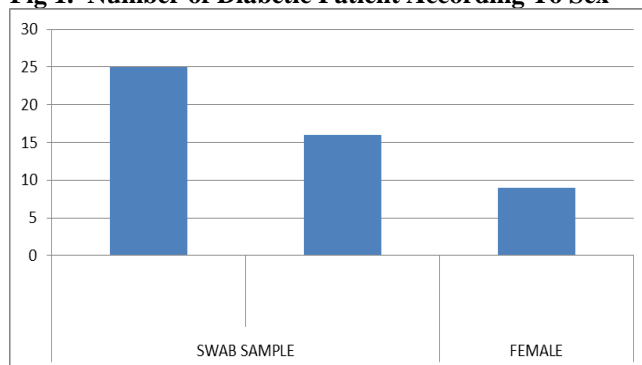
Table 1. Number of diabetic patient according to sex

Swab Sample	Male	Female
25±1.00	16±1.52	9±1.00

Values are mean ± SD of 6 individual observations.

Values are significant at $P < 0.001$.

Fig 1. Number of Diabetic Patient According To Sex



CHARACTERIZATION OF *STAPHYLO COCCUS AUREUS*

Microscope: Gram positive with more pus cells

Colony Morphology: Circular convex smooth golden yellow colony in nutrient agar Beta haemolytic colony blood yellow colouration in mannitol salt agar

Table 2. Biochemical Characterization of *Staphylococcus aureus*

Name of The Test	Result
Catalase	Positive
Oxidase	Negative
Coagulase	Positive
Indole	Negative
Methyl red	Positive
Voges Proskauer	Positive
Citrate	Positive
Urease	Positive
Gelatinase	Positive
Nitrate	Positive
Mannitol fermentation	Positive

CHARACTERIZATION OF *PSEUDOMONAS AERUGINOSA*

Microscope: Gram negative rod with more pus cells

Motility: Motile

Cultural characters: Large opaque irregular colonies with a distinctive musty or earth smell

On Nutrient agar: Green colour colony with diffused Pigmentation

On MacConkey agar: Forming NFL colonies

On Blood agar: Beta haemolytic colonies

Triple sugar iron: Alkaline butt alkaline slant

Table 3. Biochemical Characterization of *Pseudomonas aeruginosa*

Name of The Test	Result
Catalase	Positive
Oxidase	Positive
Coagulase	Not Done
Indole	Negative
Methyl red	Positive
Voges Proskauer	Negative
Citrate	Positive
Urease	Positive
Gelatinase	Positive
Nitrate	Positive

CHARACTERIZATION OF *PROTEUS MIRABILIS*

Microscope: Gram negative rod with more pus cells

Motility Test: Motile

Cultural Characters: Fishy or seminal odour

On nutrient agar: Pale colour swarming growth

On MacConkey agar: Smooth colour less separate colonies

On Blood agar: Non haemolytic colonies

Triple sugar iron: Acid butt alkaline slant gas Positive H₂S(+)

Table 4. Biochemical Characterization of *Proteus mirabilis*

Name of The Test	Result
Catalase	Positive
Oxidase	Negative
Coagulase	Not Done
Indole	Negative
Methyl red	Positive
Voges Proskauer	Negative
Citrate	Positive
Urease	Positive
Gelatinase	Positive
Nitrate	Positive
Sugar fermentation	
Glucose	+
Lactose	-
Sucrose	-
Mannitol	-

CHARACTERIZATION OF *ESCHERICHIA COLI*

Microscope: Gram negative straight rod with a few pus cells

Motility Test: Motile

Cultural Characters: The colonies are large thick grayish white moister smooth opaque or partially translucent discs

On nutrient agar: Large circular convex smooth white moist colonies

On MacConkey agar: pink colonies due to lactose fermentation

On Blood agar: Non haemolytic colonies

Triple sugar iron: Acid butt alkaline slant gas production

Table 5. Biochemical Characterization of *Escherichia coli*

Name of The Test	Result
Catalase	Positive
Oxidase	Negative
Coagulase	Not Done
Indole	Positive
Methyl red	Positive
Voges Proskauer	Negative
Citrate	Negative
Urease	Negative
Gelatinase	Positive
Nitrate	Positive
Sugar fermentation	
Glucose	+
Lactose	+
Sucrose	-
Mannitol	+

Table 6. Aerobic Bacteria Isolated From Diabetic Foot Infection

Bacterial Types	Number of The Cases	Percentage (%)
<i>Staphylococcus aureus</i>	15±2.00	60±6.00
<i>Pseudomonas aeruginosa</i>	10±1.00	36±5.40
<i>Proteus mirabilis</i>	6.0±0.50	24±4.00
<i>Escherichia coli</i>	2.0±0.45	8.0±0.50

Values are mean ± SD of 6 individual observations.

Values are significant at $P < 0.001$.

Table 7. Antibiotic Sensitivity for *Staphylococcus aureus*

Name of the antibiotic	Diameter of zone result
Ampicillin	28.0±1.50 (S)
Amikacin	20.0± 1.20 (S)
Gentamicin	20.0±1.54 (S)
Vancomycin	15.0± 0.50 (S)
Methicillin	Resistant

(S) - Sensitive (I) - Intermediate (R) - Resistance

Values are mean ± SD of 6 individual observations.

Values are significant at $P < 0.001$.

Table 8. Antibiotic Sensitivity for *Pseudomonas aeruginosa*

Name of the Antibiotic	Diameter of Zone Result
Amikacin	20.0±1.20 (S)
Gentamicin	20.0± 1.00 (S)
Ciproflaxcin	19.0± 1.10 (I)
Ceftazidime	20.0± 1.24 (S)

(S) - Sensitive (I) - Intermediate (R) - Resistance

Values are mean ± SD of 6 individual observations.

Values are significant at $P < 0.001$.

Table 9. Antibiotic Sensitivity for *Proteus mirabilis*

Name of the antibiotic	Diameter of zone result
Ampicillin	12.0± 0.50 (R)
Amoxycillin	13.0± 1.00 (R)
Cefotaxime	12.0±1.00 (R)
Ciprofloxacin	28.0± 1.52 (S)
Tetracycline	18.0± 1.45 (I)

(S) - Sensitive (I) - Intermediate (R) - Resistance

Values are mean ± SD of 6 individual observations.

Values are significant at $P < 0.001$.

Table 10. Antibiotic Sensitivity for *Escherichia Coli*

Name of the antibiotic	Diameter of zone result
Streptomycin	19.0± 1.00 (S)
Ampicillin	20.0± 1.52 (S)
Polymyxin-B	19.0± 1.00 (S)
Tetracycline	20.0± 1.15 (S)

(S) - Sensitive (I) - Intermediate (R) - Resistance

Values are mean ± SD of 6 individual observations.

Values are significant at $P < 0.001$.

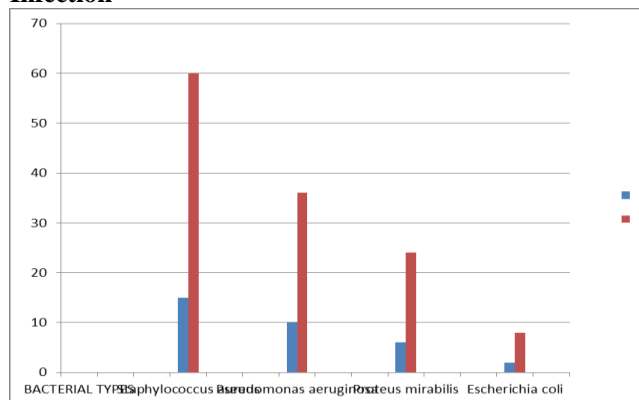
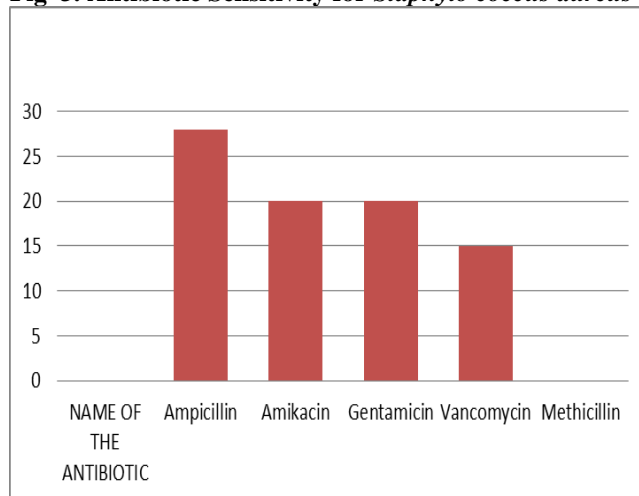
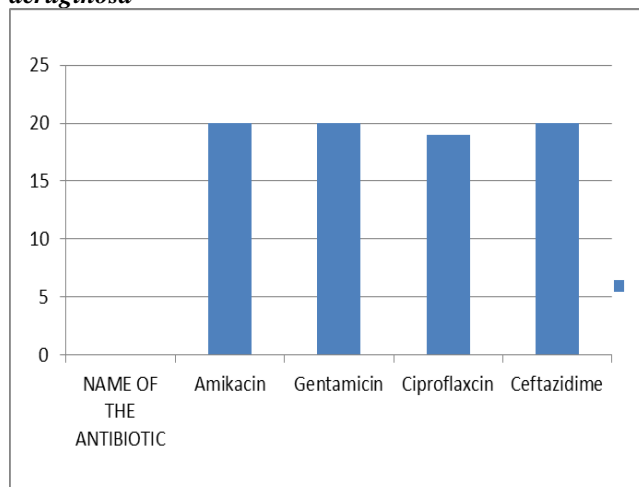
Fig 2. Aerobic Bacteria Isolated From Diabetic Foot Infection**Fig 3. Antibiotic Sensitivity for *Staphylococcus aureus*****Fig 4. Antibiotic Sensitivity for *Pseudomonas aeruginosa***

Fig 5. Antibiotic Sensitivity for *Proteus mirabilis*

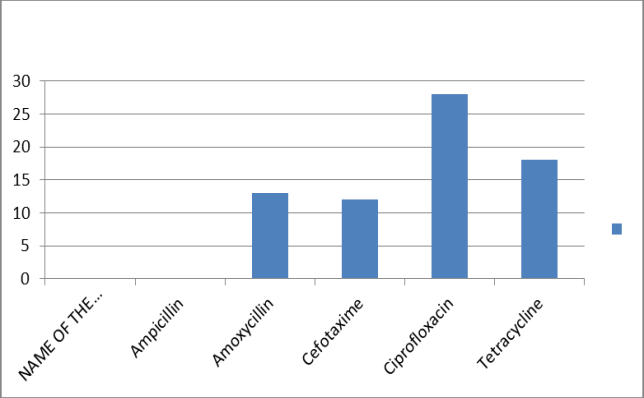
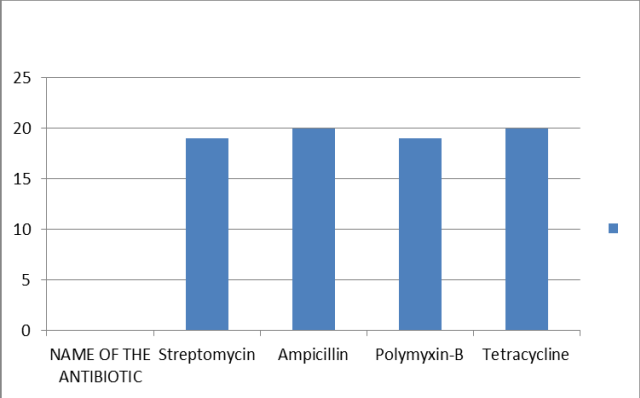


Fig 6. Antibiotic Sensitivity for *Escherichia Coli*



CONCLUSION

The human preliminary study on the diabetic foot

infected wound isolation of microorganisms and morphological arrangement in aerobic bacteria identified.

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