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Isolation and Partial Characterization of Virulent Phage Specific against Pseudomonas Aeruginosa

By Zahra Zahid Piracha, Umar Saeed, Aqsa Khurshid & Waqas Nasir Chaudhary

National University of Sciences and Technology (NUST), Pakistan

Abstract- Pseudomonas aeruginosa is an opportunistic pathogen, frequently associated with nosocomial infections worldwide. Multiple drug resistance has been reported in previous studies against pathogenic P. aeruginosa and the biofilm which makes use of antibiotics futile. Bacteriophages specific for P. aeruginosa can prove to be a new therapeutic approach for controlling infections and biolfilm contamination against this pathogen. The aim of our study was to isolate and partially characterize virulent phage specific for P. aeruginosa from sewage water. Different parameters of which make phages as suitable candidate towards future therapeutics were also investigated. Phages having lytic life cycle, high burst size and thermally more stable proposed themselves as effective therapeutic candidates. In this study, a virulent phage was isolated from sewage water having burst size of 1036 and latent period of 21 minutes. This phage has narrow host range and shows the remarkable thermal resistance and is viable up to $60 \, ^{\circ}$ C.

Keywords: pseudomonas aeruginosa, phage, isolation, partial characterization.

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Isolation and Partial Characterization of Virulent Phage Specific against *Pseudomonas* Aeruginosa

Zahra Zahid Piracha ^a, Umar Saeed ^o, Aqsa Khurshid ^e & Waqas Nasir Chaudhary ^w

Abstract-Pseudomonas aeruginosa is an opportunistic pathogen, frequently associated with nosocomial infections worldwide. Multiple drug resistance has been reported in previous studies against pathogenic P. aeruginosa and the biofilm which makes use of antibiotics futile. Bacteriophages specific for P. aeruginosa can prove to be a new therapeutic approach for controlling infections and biolfilm contamination against this pathogen. The aim of our study was to isolate and partially characterize virulent phage specific for P. aeruginosa from sewage water. Different parameters of which make phages as suitable candidate towards future therapeutics were also investigated. Phages having lytic life cycle, high burst size and thermally more stable proposed themselves as effective therapeutic candidates. In this study, a virulent phage was isolated from sewage water having burst size of 1036 and latent period of 21 minutes. This phage has narrow host range and shows the remarkable thermal resistance and is viable up to 60 °C. Sincere efforts in term of identification, isolation, purification and characterization of multiple types of phages against Pseudomonas aeruginosa; and development of cocktail with pool of lytic phages against it can prove to be a promising strategy to overcome frequent bacterial proliferation.

Keywords: pseudomonas aeruginosa, phage, isolation, partial characterization.

I. INTRODUCTION

seudomonas aeruginosa is Gram negative, rod shaped opportunistic pathogen of animals, ubiquitous in nature [1]. According to United States National Nosocomial Infection Surveillance System, it accounts for 16.1% of all nosocomial infections and ranked second among Gram negative pathogens [2]. Healthy adults rarely encounter its infection but main target is people having compromised immune system including HIV infections. The infection ranges from self-limiting folliculitis to life threatening bacteremia, wound related morbidity, septicemia, skin infections, otitis media, ecthymegangrenosum, the black necrotic lesion, endocarditis, corneal ulceration and device related infections [3-4]. It is the third leading cause of 12 % of hospital-acquired urinary tract infections, upper and lower respiratory tract infections like cystic fibrosis that is associated with high mortality rate in immune-compromised patients [4]. Gender-wise prevalence showed 61.78% male and 38.22% females were infected by *P. aeruginosa* [3]. Souli and colleagues (2008) [5] published data from 23 countries on the European Antimicrobial Resistance Surveillance System and it was shown that 18% of all isolates were multidrug resistant P. aeruginosa strains. Aman Ullah and colleagues (2012) [6] carried out a study in Islamabad and showed that P. aeruginosa is 94% resistant to Chloramphenicol, 88% to Colistin /sulphate, 73% to Tetracycline and 3% to Imipenem. The resistance against the newly tested drugs is still evolving as P. aeruginosa is highly resistant to antibiotics, both at the genetic level and as a result of living in multilayered and complex biofilm [7].

Bacteriophages are bacterial viruses that cause lethal effect on bacteria. They have genetic material in the form of either DNA or RNA (single or doublestranded), encapsidated by a protein coat [8]. Therapeutic role of the phages was interrupted by the advent of antibiotics however the emergence of multidrug resistant bacteria and adverse effects of antibiotics led to renewed interest in phages as therapeutic agent. [9]. Phages have several advantages over antibiotics and other antimicrobial agents, such as host specificity, no side effects, and multiplication in the presence of their hosts [8]. Phages are highly specific and as they are living entity so they evolve with the evolving bacteria [10]. The specificity of phages sometimes may be considered as a possible disadvantage because there are much more pathogenic bacteria than expected to be targeted. To address this problem, a cocktail of phages should be prepared [8]. The Food and Drug Administration of the United States of America recently approved some phages as safe for food products to control Listeria infections [11]. Animal tests of phage therapy are being conducted for treatments of various bacterial infections, and many lytic phages have been isolated and tested for such applications. Phages have been used to treat the Escherichia coli infections and P. aeruginosa infections in mouse models and in guinea pigs and efficient results have been obtained [12].

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Approximately 10³⁰ bacteriophages are present in the environment but only about 300 phages have been characterized [13-14]. In the pre antibiotic era, phage therapy was failed because uncharacterized phages were used so in order to use them fully, it is important to isolate and characterize new phages especially in light of the observation that most of the disease-causing organisms live in matrix-enclosed environments called biofilms that inherently show increased resistance toward all antibiotics [15-16]. In the present study, we have reported the isolation and partial characterization of a virulent phage specific for *P. aeruginosa* from sewage water in Islamabad and evaluated the different parameters of phage that makes it suitable candidate for future therapeutics.

II. MATERIALS AND METHODS

a) Identification of bacterial isolate

Pure cultures of bacterial strain were obtained from Microbiology laboratory of Kahuta research lab hospital, Islamabad. The bacteria were already resistant to Chloramphenicol, Colistin, Cotrimoxazole, Tetracycline and Aztreonam. After overnight incubation of bacterial strain, microbiological methods such as colony morphology and Gram staining were used for identification of bacterial strain [17].

b) Phage enrichment and isolation

Sewage water was taken from Sewage treatment plant I-9 Islamabad. Sewage water was centrifuged at 10,000 rpm for 10 min to remove algal cell and sewage debris. To enrich the phage population, above prepared sample concentrates (5 ml) were added to a 30 ml log phase *P. aeruginosa*. Enriched cultures were incubated overnight at 37 °C with shaking at 150 rpm. Chloroform (1%) was added to 1.5 ml of sample to disrupt bacterial cell and release phages and then centrifuged at 14,000 rpm for 10 min at 4 °C. The supernatant was filtered using 0.45 and 0.20 mm (Minisart, SalotriusStedim Biotech) syringe filters and transferred to a new tube. Phage isolation and detection was carried out by plaque assay on LB agar plate (tryptone 10 g/L, yeast extract 5 g/L, NaCl 10 g/L, agar 15 g/L) with soft agar (tryptone 10 g/L, yeast extract 5 g/L, NaCl 10 g/L, agar 0.7%) . Overnight bacterial culture and phage sample were mixed in 0.7% LB soft agar (tryptone 10 g/L, yeast extract 5 g/L, NaCl 10 g/L, agar 0.7%; at a temperature of 45 °C) and poured over LB plates. On solidification plates were incubated overnight at 37 °C and examined for plaques the following day. For a negative control, phage alone was added to the molten agar. Well isolated plaques were serially propagated until a single phage type was obtained. The purified phages were then stored in SM buffer (100 mMNaCl, 8 mM MgSO4, 50 mM, Tris-HCl [pH 7.5]), and 0.002% w/v gelatin at 4 °C with the addition of 7% dimethyl suloxide (DMSO) at -80 °C. [18]

c) One step growth experiment

One step growth curve experiment was performed to determine latent period and burst size of phage [19-20]. In brief, 50 ml of *P. aeruginosa* culture was incubated to mid exponential phase having O.D₆₀₀ 0.6 and cells were harvester via centrifugation. The pellet was re-suspended in 0.5 ml of LB media and mixed with 0.5 ml of the phage solution having plaque forming unit (pfu) of 2.75x10⁹. This mixture was allowed to stand for 3 minutes at 37 °C so that phages adsorbed to the host cells. Mixture was then centrifuged at 13,000 rpm for 2 minutes to remove the free phage particles. The pellet was re-suspended in 100 ml of LB medium and culture was incubated at 37 °C with shaking. Samples were taken after every 3 minutes for 45 minutes and after centrifugation at 13,000 rpm for 1 minute, subjected to determination of phage titer via plaque assay.

d) Analysis of calcium ion effect on phage adsorption

In order to measure the effect of divalent metal ions on phage adsorption rate $CaCl_2$ was used [21]. The 50 ml of *P. aeruginosa* overnight culture having $O.D_{600}$ 0.6 was divided in two autoclaved flask 25 ml each. One flask was inoculated with phage 500 µl having pfu of 2.75x10⁹ (control), while the second flask was inoculated both with 500 µl phage (pfu 2.75x10⁹) and 250 µl of 1 M CaCl_s and incubated with constant shaking at 90 rpm at 37 °C. Samples were taken from both flasks at different time intervals of 0, 10, 20, and 30 minutes. Samples were centrifuged at 13,000 rpm for 3 minutes to sediment the phage adsorbed bacteria. The supernatant was assayed for unabsorbed phages via double layer plaque assay method and counts were compared with the titer of control.

e) Thermal stability analysis

Thermal stability tests were performed according to the method described by Suarez [21]. Phage filtrates were taken in microcentrifuged tubes and treated under different temperatures at 37 °C (control), 40 °C, 50 °C, 60 °C and 70 °C for 20 minutes, 40 minutes and for 1 hour. After incubation plaque assay was performed for each treated sample. Results were compared with control.

f) Bacterial reduction assay

P. aeruginosa culture 1 ml was inoculated in 2 flasks having 150 ml LB media. The phage supernatant 500 μ l was introduced in one flask and other flask was treated as control and both the flasks were incubated in shaking incubator at 37 °C. The O.D₆₀₀ readings were taken after every 2 hours for 24 hours using spectrophotometer. The values were compared with control.

g) Host range

The host range of the isolated phage was checked on the range of clinical pathogenic bacterial

strains of *P. aeruginosa* isolated from blood, wound, urinary tract, ear and pus, *Streptococcus pneumoniae, Escherichia coli* and *Klebsiella pneumoniae*. All the tested bacterial strains were clinical pathogens obtained from Microbiology laboratory of Kahuta research lab hospital, Islamabad. Susceptibility of the phage was tested via the spot on lawn technique [22]. Plates were incubated upside down overnight and checked for any plaque against the negative control i.e., uninfected lawn.

III. RESULTS

a) Identification of bacterial isolate

In this study, *P. aeruginosa* was identified via Gram staining and colony morphology. Gram staining showed that it is Gram negative bacillus.

Table 1: Colony morphology characteristics of P. aeruginosa							
Sr.	Color	Margin	Shape	Elevation	Luminous	Texture	Size
INO					benavior		
1.	Off	Undulat	Irregula	Raised,	Translucen	Mucoid	1
	white	е	r	spreading	t		mm
				edges			

b) Bacteriophage isolation

P. aeroginosa was used as indicator strain for phage isolation. After enrichment, phage containing samples were placed on semisolid LB medium with *P. aeruginosa* forming lawn. Two phages were isolated that ranged from 3 to 4 mm in diameter on lawn of indicator strain by plaque analysis (Fig 2). These phages were isolated from sewage water, the same reservoir as their host. Later experiments showed that the observed plaques of two phages were similar in all aspects. Purified phage produced clear plaques on lawn of host bacterium *P. aeruginosa* (Fig 3).



Figure 2: Master plate of plaque assay for phage isolation from sewage water against *Pseudomonas aeruginosa*, marked points show the presence of bacterial growth clearance in the form of round zones (plaque)



Figure 3: Double layer agar plates showing plaque purified after repeating plaque assay three times and filtering phage lysate through 0.2 mm filter. Calculated PFU/ml was 2.75x10⁹ and observed plaque size was 3-4 mm

c) One step growth experiment

One step growth curve experiment was performed to determine the latent period and burst size of the phage. Latent period is the interval between adsorption of phages to host cell and cell lysis and burst size is progeny phage particles produces by a single host bacterium was determined [19-20]. The results were analyzed and the number of infective centers or constant phage titer along with the latent stage was deduced. A tri obtained including the latent phase, rise or log phase and stationery or plateau phase. The latent time of the phage was determined to be 21 minutes and burst size of the phage was 1036 phage/ infected cell. It was based on ratio of mean yield of phage particles



Figure 1 : Gram staining of *Pseudomonas aeruginosa* showing Gram negative bacillus rods

liberated to mean phage particle that infected the bacterial cells. There was no difference in burst size

between a 1-month-old phage stock and a freshly prepared phage stock.



Figure 3: One step growth curve experiment showing the latent period (21 min) and the average burst size (1036 viral particles per host cell). Latent time and burst size of phage were inferred from the curve with a tri phasic pattern. L, latent phase; R, rise phase; P, plateau phase

 d) Calcium Ion Effect on the Adsorption Rate of Phage Divalent ions often affect the adsorption of phages. In order to check this, CaCl₂ 1mM was added in *P. aeruginosa* phage mixture and number of free phages left in solution were detected after 10 minutes interval for 30 minutes by using plaque assay. Data analysis showed that significant differences existed between the control and calcium ion treated group. Results obtained show that calcium ions stabilize the process of adsorption and infectivity is increased due to addition of CaCl₂.



Figure 4: Adsorption rate of phage at different time intervals shows that divalent metal ions effect on adsorption rate

e) Thermal stability analysis

Thermal stability test was carried out to check the infectivity of phage at different temperatures at pH 7.0 and also to check the resistant capability of phages. The preliminary experiments show that phages remain active at 4 °C for about 4 months (data not shown). So stability at higher temperatures was checked. Phages were amazingly stable at 35 °C, 40 °C, 50 °C and 60 °C for 20, 40 and 60 minutes. Results showed that phages are heat resistant and even 16 % phages remain viable up to 60 °C after 1 hour incubation.



Figure 5: Thermal stability of phages was analyzed at different temperatures and surviving phages were tittered via plaque assay

f) Bacterial reduction assay

Infection of bacteria with the phages was monitored for 24 hours. The bacterial reduction caused by phages was compared with control. Phage infection produced a drastic decrease of the *P. aeruginosa* culture as compared to control as it is evident from the graph. However, a constant increase in OD₆₀₀ was seen after 16 hours this is most probably due to growth resistant phages.





g) Host range

The susceptibility of the phage was investigated with clinical strain of *P. aeruginosa* isolated from various

sources and results show that the phage hasnarrow host range.

Sr. No	Source of bacteria	Clinical strain	Host range (+,-)
1.	Pus sample	P. aeruginosa	+
2.	Blood sample	P. aeruginosa	-
3.	Urinary tract sample	P. aeruginosa	_

Table 2: Host range determination of the	phage

4.	Wound sample	P. aeruginosa	+
5.	Ear sample	P. aeruginosa	+
6.	Ear sample	S. pneumoniae	-
7.	Urinary tract sample	E. coli	-
8.	Urinary tract sample	K. pneumoniae	-

IV. DISCUSSION

P. aeruginosa is a nosocomial pathogen and leading cause of health care associated infections. It is an emerging multidrug resistant pathogen around the globe and also in Asia [2]. Way of acquiring resistance specifically the chromosome encoded efflux and low outer membrane permeability accounts for its high resistance and makes use of antibiotics futile. The biofilm induces the resistance and restricts the diffusion of the antibiotic which makes most of the antimicrobial agents less effective. It can also acquire additional resistance genes from other organisms via plasmids, transposons and bacteriophages [7]. Imipenem was considered one of the effective antibiotics however Lautenbech and colleagues in 2010 detected the imipenem resistant P. aeruginosa strains. All these evidence showed an alarming situation and demands an alternative treatment for *P. aeruginosa* infections [6, 23].

In this study, our results showed that a newly isolated bacteriophage can lyse actively growing *P. aeruginosa.* Phages are found with the natural habitat of their host cells such as sea water, sewage water/ sludge ponds etc [1]. They are host specific and evolve along with their host [10]. Phage was isolated from sewage water. Sewage in general, contains a large diversity of microbes due to contamination from fecal and hospital drainage water. This newly isolated phage was highly lytic and produced large plaque of 4 mm.

Phage establishes an infection in the host when it gains entry into the host after interaction with the bacterial receptor [24]. Many phages have been found to be greatly specific for their receptors present on the host cell surface. They show no interaction with receptors having a different structure. The specificity turns out to be the basis of phage typing methods used for the identification of bacterial strains. The results obtained clearly showed that our virulent phage for *P. aeruginosa* was highly specific. Results indicated that the phage may be using a common receptor.

One step growth curve explains all the stages involved in multiplication of phages. From this curve, latent period and burst size are determined. Latent period or incubation period is the time which the phage requires inside the bacterial cell and is 21 minutes in case of our isolated phage. The optimum latent period leads to high phage fitness. Rise phase is the time when infected cell begin to lyse and plateau phase is time when the infectious centers is due to only single virion [19]. Phage burst size is average number of phage bacterial cells produced per bacterial cells upon infection. The burst size of phage is 1036 particles per bacterial cell. Wang [25] described that the duration of the latent phase correlates with the burst size. The longer is the latent phage the bigger is the burst size of the phage. The number of phages liberated upon infection is quite high as compared to early reported phages.

The factors which influence the infectivity of phage in-vitro are temperature and divalent cations. The physiological state of host also plays an important role in reproduction of phage. The infection of the virion starts when it binds to host bacterial cell. In phage host interaction, calcium might be playing an important role in infection cycle [19] .Calcium ion stabilizes the weak interaction of virion with receptors or they may change the structure of the receptor to make the accessibility of phages smoothly. Ions have an electrostatic effect in the interaction of phage bacterium systems [26]. Different amount of calcium gives different infectivity.

The physical parameters like the temperature not only accelerate the adsorption of phages but also have direct effect on metabolic activities of the host [26]. Extremely thermal resistant phages have been isolated from thermal habitats and also in other aquatic environments [27]. Our results implied that phage we have isolated is active on high temperatures. The phage remained active till four months at 4 °C and it remains viable up to 60 °C after 1 hour incubation. The maximum infectivity was observed at 37 °C and least infectivity was observed at 60 °C.

The phage we isolated also causes decrease in turbidity of *P. aeruginosa* culture and this behavior was obvious for 16 hours as compared to control. There was maximum bacterial destruction during this time period but some bacterial cells show resistance to virus infection. This may be because of host bacterium adaptation as bacterial replication was going on meanwhile and bacterial resistant cells survived and though they were less in number but they started multiplying after about 16 hours and it was time when turbidity of culture increases as shown in figure 6. This type of behavior can be a hindrance in phage therapy. Some studies shown that phage resistant bacteria losses their virulence factor because those virulence can be a site for phage infections. Such loss of virulence factor in a phage resistant bacteria mutant has been demonstrated in fish pathogens [28]. Due to resistance, the fitness of bacteria reduces and it renders it competing unfavorably with its phage sensitive ancestors. The gradual increase in optical density can also be attributed to some other factors like bacterial debris and this might be a hindrance in exposition of actual results.

V. Conclusion

Viral diseases are increasing day by day in world and it is anticipated that soon hepatitis viruses would emerge as most dangerous viral pathogens [29,30]. But on the other hand nature has selected many viruses to kill various other pathogens. . Nature has many hidden remedies against multiple disorders such as cardiovascular diseases, metabolic disorders, cancer, chronic inflammation and many others); there is strong need to discover therapeutic potentials of natural items. Many studies showed elevated activation of various cellular proteins cause cancer proliferation, which can be further inhibited by potential inhibitors (31-35). Partial characterization of the virulent phage showed that it is highly efficient in lysing Pseudomonas aeroginosa, as it has shown some outstanding aspects including rapid growth nature, high thermal stability and optimum latent period. All these characteristics make this phage very promising for possible application in eradication of Pseudomonas aeroginosa contaminations and treatment of Pseudomonas aeroginosa infections. Phage has shown narrow host range, so for the broad-spectrum elimination of bacteria; a "cocktail" with a pool of lytic phages might be more useful against present and other bacterial strains. A better understanding of phages and lytic enzyme biology could facilitate development of novel future therapeutics against multiple drug resistant bacteria.

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Degradation of Dimethoate by Cellulolytic Bacteria in Cotton Soils

By Dr. R. Jaya Madhuri

Abstract- The present experience has been aimed to isolate bacteria specifically cellulolytic bacteria, normally found in cotton soils and determine their role in degradation of the specific pesticide. Two cellulolytic bacterial isolates k1 & k2 identified as Pseudomonas putida and Bacillus pumulus were capable of growing on dimethoate supplemented medium. P.putida exhibited maximum growth of 4.1x 106 cfu/ ml at 0.09 mg/ml while B.pumulus showed significant growth of 2.2 x107 cfu/ml at 0.06mg/ml of dimethoate after 72hrs of incubation at room temperature. Rate of utilization of dimethoate increased progressively with increase in the concentration of yeast extract added to the medium up to 0.56% in P.putida and 0.7% in B.pumulus. The present findings indicate that among the two bacterial isolates, sps of Pseudomonas could degrade 88% of dimethoate while Bacillus sps exhibited high degradation rate of 92% which can be commercialized for bioremediation of dimethoate contaminated sites. Further, the rate of degradation is maximum at 72hrs of incubation.

Keywords: dimethoate, cellulolytic bacteria, degradation.

GJMR-C Classification : NLMC Code: QV 350

DE GRADATIONOFDIMETHOATE BY CELLULOLYTIC BACTERIAINCOTTONSOILS

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Abstract- The present experience has been aimed to isolate bacteria specifically cellulolytic bacteria, normally found in cotton soils and determine their role in degradation of the specific pesticide. Two cellulolytic bacterial isolates k1 & k2 identified as Pseudomonas putida and Bacillus pumulus were capable of growing on dimethoate supplemented medium. P.putida exhibited maximum growth of 4.1x 106 cfu/ ml at 0.09 mg/ml while B.pumulus showed significant growth of 2.2 x107 cfu/ml at 0.06mg/ml of dimethoate after 72hrs of incubation at room temperature. Rate of utilization of dimethoate increased progressively with increase in the concentration of yeast extract added to the medium up to 0.56% in P.putida and 0.7% in B.pumulus. The present findings indicate that among the two bacterial isolates, sps of Pseudomonas could degrade 88% of dimethoate while Bacillus sps exhibited high degradation rate of 92% which can be commercialized for bioremediation of dimethoate contaminated sites. Further, the rate of degradation is maximum at 72hrs of incubation.

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I. INTRODUCTION

otton, the most important fibre crop of India plays a dominant role in its agrarian and industrial economy. It is attacked by various sucking pestsjassids, aphids, thrips, whitefly, red spider, mite and mealy bug. Bollworms such as pink bollworm, spotted bollworm and American bollworm (Tomelin, 1995).

The progressive increase of pest problem and demand for agricultural products necessitated the application of agrochemicals and ensure high quality and crop yield (Graebing et.al.,2002). Among the xenobiotics used, organophosphorous are widely applied in Indian agricultural system.

The introduction of these pesticide in to the soil environment raises concern as to their effect on ecological balance in terms of soil fertility (Balwinder Singh 2002). Even though the pesticide degradation is under the impact of various physico-chemical parameters, literature survey reveals that the major pathway governing degradation and ecotoxicity of these compounds is *microbial* mediated (Latha et al, 2001).

Dimethoate, an organophosphate broad spectrum insecticide is of particular concern as it is widely used for controlling cotton pests. But in view of its toxicity, it is important to remove dimethoate from the environment. The role of microorganisms in bioremediation is important because of their ability to degrade hazardous compounds into harmless ones. There is lack of information regarding biodegradation of dimethoate in cotton soils.

Hence the present work has been aimed to characterize the isolated cellulolytic bacteria from the cotton soil and for its ability to utilize dimethoate as the sole source of carbon and energy.

II. MATERIALS AND METHODS

For the isolation of bacteria, soil samples from cotton cultivated fields were collected from three different locations in Guntur District (Narsarao pet, Sattena Palli and LAM farm). All soil samples were sieved through a 2mm screen and then used.

a) Isolation and characterization of cellulolytic bacteria

The three soil samples were enriched with plant debris and incubated with 37° C for 15 days. After the incubation period, enriched soil samples were taken and serial dilutions (up to 10^{-6}) were made using sterile water. One ml aliquots of 10^{-4} and 10^{-5} dilutions were made to the sterile cellulose agar medium allowed to solidify. Triplicates for each dilution were maintained.

The plates were incubated at 37°C for 3-5 days and were flooded with 1% aqueous solution of hexadecyltrimethyl ammonium bromide, the isolates showing a clear zone of hydrolysis around the colony were selected for further study. These isolates were streaked on cellulose agar slants and maintained as pure cultures.

Biochemical characterization of isolated bacteria was done referring to the Bergey's manual of Systematic Bacteriology and named the two bacterial isolates producing maximum zone of hydrolysis as K1 and K2

b) Utilization of Dimethoate

The rate of dimethoate utilization in terms of growth by the isolates K1 and K2 was determined at 37° C in dimethoate (0.1 mg/ml) mixed mineral salt solution for 140 hrs.

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c) Dimethoate degradation test

The degradation of dimethoate was observed by bioassay method. 25 ml of MS solution was taken in 100 ml conical flask and 0.03, 0.06, 0.09, 0.12, 0.15, 0.18, 0.21 and 0.24 mg/ml of dimethoate were taken to which the bacterial inoculum of K1 and K2 separately were added and incubated at 37°C for 72 hrs and growth was studied by colony count technique.

d) Effect of yeast extract on the utilization of dimethoate

The effect of varying levels of yeast extract on the rate of dimethoate utilization was studied for the isolated bacteria K1 and K2. The bacterial isolates were allowed to grow for 24 hrs at 37° C in MS solution amended with dimethoate(0.1 mg/ml) plus yeast extract (MSDY) and in MS solution supplemented with yeast extract(MSY) only as control. The results were tabulated and interpreted in terms of viable CFU/ml in MS solution.

Dimethoate degradation test : Bacterial cultures efficient in degradation of dimethoate were incubated for 72 hrs and the percentage of degradation as well as metabolites formed are analysed by HPLC technique using hexane as mobile phase and C18 column at IIHR, Bangalore.

III. Results

a) Isolation and characterization

The K1 isolate obtained from cotton soil enriched with plant debris is a gram negative motile rod with 2.2 μ m length and 1.3 μ m width. It is aerobic and showed no spore formation. It is non spore forming, motile, catalase, oxidase and casein positive. Starch and gelatin are not hydrolyzed. IMVIC negative and nitrate is not reduced. According to Bergey's manual of classification, the K1 isolate was tentatively identified as *Pseudomonas putida* (Table 1).

The K2 isolate obtained is a gram positive rod with 3μ m length and 0.6μ m width. It is aerobic and a terminal ellipsoidal spore was observed. Utilization of urea, gelatin, citrate and nitrate reduction was observed. Hydrolysis of casein and starch was observed. The isolate was catalase positive and oxidase negative. Based on the above characteristics the K2 isolate was identified tentatively as *Bacillus pumilis*.

Table 1 : Morphological and biochemical
characterization of K1 and K2

Name of the reaction	K1	K2
Gram reaction	Negative	Positive
Morphology	Rod	Rod
Cell length	2.2 µm	3 <i>µ</i> m
Cell diameter	1.3 <i>µ</i> m	μm
Pigment	Yellow	No pigmentation

Spore formation	Absent	Terminal, Ellipsoidal
Motility	Motile	Nonmotile
Starch hydrolysis	Negative	Positive
Catalase test	Positive	Positive
Gelatin hydrolysis	Negative	Positive
Oxidase test	Positive	Negative
Urease test	Negative	Positive
Casein hydrolysis	Positive	Positive
Nitrate reduction	Negative	Positive
Indole production	Negative	Negative
Voges Proskaeur	Negative	Negative
Citrate utilization	Negative	Positive

b) Utilization of Dimethoate

The growth curve of K1 and K2 isolates cultivated in 0.1 mg/ml of dimethoate is shown in tables 2 and 3. The K1 isolate showed its maximum growth rate after 20 hrs in MS solution (Tab 2).

The K2 isolate showed its maximum growth rate after 40 hrs in MS solution (Tab 3).

Table 2 : Growth of k1 in mineral salt solution
supplemented with 0.1mg/ml of dimethoate

Hours	CFU/ml
0	1.2x 10⁵
20	3 x 10⁵
40	4.5 x 10⁵
60	1.2 x 10⁵
80	4 x 10 ⁴
100	8 x 10 ³
120	6 x 10 ³

Table 3 : Growth of k ₂ in mineral salt solution
supplemented with 0.1mg/ml of dimethoate

Hours	CFU/ml
0	1.5 x 10⁵
20	2.3 x 10⁵
40	3.8 x 10⁵
60	1.2 x 10⁵
80	4.1 x 10 ⁴
100	5.8 x 10 ³
120	3.4 x 10 ³

c) Dimethoate tolerance and utilization test

Degradation of dimethoate provided as the sole carbon was studied in mineral salt solution. In K1 isolate it was found that maximum growth occurred at 0.09 mg/ml after incubation for 72 hrs at 37°C (Tab. 4). The K2 isolate was found tolerating dimethoate up to 0.24mg/ml in MS solution with maximum growth at 0.06mg/ml after an incubation for 72hrs at 37°C (Tab 5).

Table 4 : Utilization and tolerance level of dimethoate by $$k_1$$ isolate

Dimethoate concentration (mg/ml)	CFU/ml
0	1.5 x 10 ⁴
0.03	2 x 10 ⁴
0.06	2.7x 10 ⁴
0.09	4.1 x 10 ⁶
0.12	2.4 x 10 ⁴
0.15	1.3 x 10 ³
0.18	8.8 x 10 ²
0.21	No growth
0.24	No growth

Table 5 : Utilization and tolerance level of dimethoate) by
k ₂ isolate	

Dimethoate concentration (mg/ml)	CFU/ml
0	1.3 x 10 4
0.03	1.9 x 10 ⁴
0.06	2.2 x 10 ⁷
0.09	1.1 x 10⁵
0.12	2.3 x 10 ⁴
0.15	1.8 x 10 ³
0.18	1.4 x 10 ³
0.21	0.8 x 10 ²
0.24	0.2 x 10 ²

d) Effect of Yeast Extract on the Utilization of Dimethoate

The K1 isolate when used for the utilization of dimethoate in the presence of yeast extract showed higher degrading activity. The rate of utilization of dimethoate increased progressively with the increase in concentration of yeast extract up to 0.56% but the growth was effected at a concentration of 0.7% and above (Tab 6).

The K2 isolate when used for the utilization of dimethoate in the presence of yeast extract showed higher degrading activity. The rate of utilization of dimethoate increased progressively with the increase in concentration of yeast extract up to 0.70% but the growth was drastically effected at a concentration of 0.84% and above (Tab 7).

Table 6 : Effect of yeast extract on k1 utilization ofdimethoate (0.1 mg/ ml) in mineral salts solution withinitial inoculum of 1 x 103 CFU/ml

Conc. of yeast extract	Mineral salts medium with dimethoate	Mineral salts medium with dimethoate and yeast extract
0.14	2.1 x 10⁴	2.9 x 10 ⁴
0.28	2.8 x 104	3.5 x 10⁵
0.42	3.5 x 10⁵	1.7 x 10 ⁶
0.56	4.7 x 10⁵	4 x 10 ⁶
0.70	3.3 x 10 ⁴	3.1 x 10⁵
0.84	2.5 x 10 ⁴	5.1 x 10 ⁴
0.98	1.6 x 10 ³	2.8 x 10 ⁴

Table 7 : Effect of yeast extract on k_2 utilization ofdimethoate (0.1 mg/ ml) in mineral salts solution withinitial inoculum of 1 x 103 CFU/ml

Conc. of yeast extract	Mineral salts medium with dimethoate	Mineral salts medium with dimethoate and yeast extract
0.14	1.5 x 10 ⁴	1.9 x 10 ⁴
0.28	1.9 x 10 ⁴	2.5 x 10⁵
0.42	3.1x 10⁴	1.5 x 10 ⁷
0.56	5 x 104	3.9 x 10 ⁷
0.70	1 x 10 ⁵	3.1 x 10 ⁸
0.84	1.5 x 10⁵	7.9 x 10⁵
0.98	2.5 x 10⁵	3.9 x 10 ⁴

Table 8 : Rate of degradation of dimethoate by Bacilluspumulis K_1 isolate

Incubation time	% of degradation
24 hr	38
48 hr	51
72hr	88

 Table 9 : Rate of degradation of dimethoate by Bacillus pumulis K2 isolate

Incubation time	% of
	degradation
24 hr	43
48 hr	78
72 hr	92

e) Degradation of dimethoate by selected bacterial cultures

Bacterial cultures obtained by enrichment technique were tested for dimethoate degradation. In the present experiment, at the end of incubation, HPLC analysis shows that 38% degradation occurred at 24 hrs. At 48 hrs time interval, 51% dimethoate disappeared. Finally after 72 hrs,only 12% of the parent residue was detected in the sample. That means 88% of compound was degraded with respect to k_1 . (Tab 8)Regarding k_2 , only 85% of the parent compound was detected at the end of incubation period. Even though previous evidence indicate rapid degradation of the pesticide by soil bacteria, more than 90% of degradation rate was not reported. But in the present experiment, only trace amount of the pesticide residue was remaining after bacterial degradation indicating high efficiency of K₂ (. Morover, formation of extra peaks besides the parent compound indicate formation of intermediate products means biotransformation of the which parent insecticide. But due to non availability of compound, end product is not identified.

IV. DISCUSSION

In the present study, two bacterial isolates K1 and K2 which are cellulolytic and capable of growth on dimethoate as a sole source of carbon and energy have been isolated from the cotton soils.

In this study the strain K1 showed maximum growth of 4.1x 10⁶cfu/ml at 0.09mg/ml and strain K2 with significant growth of 2.2x10⁷cfu/ml at 0.06 mg/ml. Bhadbade et al, (2002) reported that Bacillus lichen-iformis tolerated dimethoate upto the concen-tration of 2 mg/ml.

The K1 isolate showed maximum utilization of dimethoate at a concentration of 0.09 mg/ml and K2 isolate showed maximum utilization of dimethoate at a concentration of 0.06 mg/ml after 72 hrs at 37°C. In a previous study Mandal et.al (2002) reported that *Bacillus licheniformis* showed a maximum utilization of dimethoate at 0.45 mg/ml

The K1 isolate when used for the utilization of dimethoate in the presence of yeast extract showed higher degrading activity. The rate of utilization of dimethoate increased progressively with the increase in concentration of yeast extract up to 0.56% but the growth was effected at a concentration of 0.7% and above.

The K2 isolate when used for the utilization of dimethoate in the presence of yeast extract showed higher degrading activity. The rate of utilization of dimethoate increased progressively with the increase in concentration of yeast extract up to 0.70% but the growth was drastically effected at a concentration of 0.84% and above. Sharmila et al, (1992a and b) also reported earlier that rate of microbial degradation of parathion and other organophosphates in soil was regulated by the amount of yeast extract in the medium. Similar evidence was provided by Despande et al, (2001). Accordingly, rate of utilization of dimethoate by Bacillus sps progressively increased with yeast extract concentration up to 0.15%. This was further supported experimental results of Kadam (2003) at bv Sreerampore. India.

Even though previous evidence indicate rapid degradation of the pesticide by soil bacteria, (Dixit and Banerjee, 2000; Fernandez etal, 2002; Zabik etal;2003).more than 90% of degradation rate was not reported. But in the present experiment, only trace amount of the pesticide residue was remaining after bacterial degradation indicating high efficiency of K_2 .

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Diagnostic Role of the Bethesda System for Reporting Thyroid Lesions: Effective Tool for Managing Thyroid Lesions

By Dr. Vasudha M. Bhagat, Dr. Hemali J. Tailor, Dr. KumarBhargav R. Kaptan, Dr. Varsha Baladawa, Dr. Gunjan H. Prasad, & Dr. Peeyush K. Saini

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Abstract- Introduction: As Fine needle aspiration cytology (FNAC) is the primary investigation for the management of thyroid lesions, its interpretation is very crucial. The Bethesda System for Reporting Thyroid Cytopathology (TBSRTC) for clarity of communication recommends that each case should be reported in 1 of 6 general diagnostic categories facilitating communication among the managing team of doctors and leaves almost no confusion regarding the management of thyroid lesions.

Aim: To study utility of The Bethesda system of reporting thyroid cytopathology.

Materials and Method: During period of 1 year from 1st January to 31st December 2012 aspiration cytology has been carried out in 160 thyroid swellings referred to cytology department of a tertiary care hospital in Surat. Fine needle aspiration cytology was performed using mainly non-aspiration and aspiration techniques. All the cases were reported using TBSRTC. Cases were followed whenever possible.

Keywords: FNAC, non aspiration technique, thyroid lesions, the bethesda.

GJMR-C Classification : NLMC Code: WK 200

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Diagnostic Role of the Bethesda System for Reporting Thyroid Lesions: Effective Tool for Managing Thyroid Lesions

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Abstract- Introduction: As Fine needle aspiration cytology (FNAC) is the primary investigation for the management of thyroid lesions, its interpretation is very crucial. The Bethesda System for Reporting Thyroid Cytopathology (TBSRTC) for clarity of communication recommends that each case should be reported in 1 of 6 general diagnostic categories facilitating communication among the managing team of doctors and leaves almost no confusion regarding the management of thyroid lesions.

Aim: To study utility of The Bethesda system of reporting thyroid cytopathology.

Materials and Method: During period of 1 year from 1st January to 31st December 2012 aspiration cytology has been carried out in 160 thyroid swellings referred to cytology department of a tertiary care hospital in Surat. Fine needle aspiration cytology was performed using mainly non-aspiration and aspiration techniques. All the cases were reported using TBSRTC. Cases were followed whenever possible.

Observations and Results: The cytological samples were assessed by qualified consultant pathologists and were categorized in category 1 to 6,six tier system according to TBSRTC criteria given by National Cancer Institute USA. Fine needle aspiration cytology analysis revealed 149 (93.12%) non-neoplastic and 11(6.88%)neoplastic lesions. Major bulks of 140 cases (87.5%) were of category II.

Conclusion: Application of TBSRTC bridges the gap in communication amongst the cytopathologist, endocrinologists, surgeons, radiologists, and other health care providers not only in the confined region but also worldwide and leaves no confusion regarding management of thyroid lesions. Few of the borderline lesions often create the confusion which are eliminated by TBSRT.

Keywords: FNAC, non aspiration technique, thyroid lesions, the bethesda.

I. INTRODUCTION

hyroid enlargement is a common occurrence in most regions of the world including India. Being tertiary care hospital we frequently encounter such cases because southernGujarat and surrounding mountainous areas are one of the endemic goitrebelt in India.

Thyroid lesions are one of the most common lesions subjected to the cytopathology as Fine needle aspiration cytology (FNAC) is the first line investigation apart from other investigations like ultrasonography (USG), thyroid function tests, thyroid scan, and antibody levels are done subsequently to select the patients who require surgery and those that can be managed conservatively.^[1, 2]

For the primaryevaluation of patients FNAC has proven to be a rapid, cost-effective, safe and reliable method of investigationlikein lesions of breast, lymph nodes and others.^[3, 4] However, the success of FNAC is dependent on several important contributing influences including aspirator experience, skilful interpretation rational analysis and its application in management. Data from the Surveillance Epidemiology and End Results (SEER) registry show an increasing prevalence of differentiated thyroid cancer worldwide, ^[5, 6]

The increasing prevalence of thyroid cancer and improvements in the technology and resolution of ultrasound machines have led to an increasing number of cytological diagnostic procedures. [7] So being the primary investigation, interpretation and application of FNAC findings is very crucial for further management of thyroid lesions especially for ruling out need of surgery. Uniform communication amongst the cytopathologist, surgeons, endocrinologists, radiologists, and other health care providers will eliminate confusion regarding management. Few borderline thyroid lesions often create confusions regarding treatment. To eliminate such dilemma National Cancer Institute USA in 2007 conference meet was organised in Bethesda with one of the objectives beingto standardize the diagnostic terminology for the reporting of thyroid cytopathology results. The recommendations resulting from this conferenceled to the formation of The Bethesda System for ReportingThyroid Cytopathology (TBSRTC). This classificationscheme has achieved its purpose of standardization of thyroid-reporting cytopathology, as evidenced by several publications.^[8]

It has been shown that the best way toimplement TBSRTC in a pathology laboratory and to

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convince the clinicians of its validity and usefulness is to compare the outcome data prior to and after the implementation of TBSRTC. Current study was mainly focussed to study the role of reporting system.

II. MATERIAL AND METHOD

Study includes 160 cases of thyroid swellings patients referred to the cytopathology section of pathology department in a tertiary care hospital in southern gujarat between January 2012 and December 2012. Patient's details regarding history, clinical examination, thyroid function tests, clinical diagnosis, FNAC and histological data whenever possible were noted. The data were analysed in simple statistical tables.

All the cases of thyroid swelling subjected to FNAC were performed by cytopathologist. Prior to procedure, palpation was carried out to note the mobility of the thyroid during swallowing and the presence of any enlarged cervical lymph node. The patients were made to lie supine with their necks stretched up. A 23-24 gauge needle was used, with non-aspiration technique in most cases and very few cases with aspiration technique by a 10 ml disposable syringe. Two or more passes at different sites were made in each case. No major complications like penetration injury to the trachea, laryngeal nerve, or hematoma were recorded. Slides were prepared from aspirated material. In the case of cystic nodules, the cysts' contents wereaspirated, centrifuged, and slides made from the sediment for cytological examination. The slides were stained with MayGrunwald Giemsa (MGG), Papanicolou [PAP] stain and Haematoxylin and Eosin (H&E) and examined under light microscope. The microscopic diagnosis was interpreted under guidelines laid down by TBSRTC including categories I to VI (table 1) after taking into account of all available clinical, radiology and other data. Whenever possible, further sub typing was given. The cytological diagnosis were correlated with clinical features, thyroid function tests, subjected to histopathological examination whenever possible.

III. **Results**

Study includes 160cases with age rangebetween 5 to 70 years. Maximum cases were in 21 – 50 years of age group. Bulk of the cases were females comprising of 136 cases (85%) and 24 cases (15%) weremales and female: male ratio was 5.67:1. Long standing history of thyroid swelling was the main presenting symptom. Swelling was mainly diffuse and nodular in few cases. Symptoms like pain in the neck region, dysphagia, hoarseness of voice and cough were rare.

FNAC of 160 patients yielded the following diagnosis as depicted in Table 1.

a) Non neoplastic

Category I: The totals of 9 patients (5.63%) were diagnosed under nondiagnostic or unsatisfactory category.

Category II: It included most of the study cases with 140 cases (87.5%) of total. Age ranges with maximum number of cases were in 20-50 year age group. It consists of 'non-neoplastic' or 'negative for malignancy' cases like colloid goitre with 97 cases (69.29%), Thyroiditis with 36 cases (25.71%) and Adenomatoid goitre 7 cases (5%). All of these benign cases were just followed up and surgery was prevented.

Category III: It includes lesions which were not clearly benign or malignant. Conclusive opinion was not possible. We did not have any case diagnosed in this category.

Category IV: It includes 5 cases(3.12%) of follicular neoplasm (FN) and suspicious of follicular neoplasm (SFN).

Category V: It includes only 1 case (0.63%).

Category VI: It included 5 cases (3.12%) comprising of 2 cases each of papillary thyroid carcinomas and anaplastic carcinoma of thyroid with 1 case of medullary carcinoma of thyroid.

IV. Discussion

As in management of thyroid lesions, FNAC is the gold standard and primary investigation of choice along with other investigations like (USG) ultrasonography examination, thyroid function tests, thyroid scan, and antibody levels are done subsequently to find out patients who require surgery and those that can be managed conservatively. ^[1, 2]

Being a tertiary care hospital we have many patients of thyroid disorders from the South Gujarat region including Bharuch, Vapi, Songadh, Vyara and other goitre belts. Majority of these lesions are usually benign and require no aggressive treatment. So interpretation in each case is very crucial for further management Also we want to establish uniform communication between the pathologist, radiologist, endocrinologist, surgeons and treating physicians. So that there would be no confusion regarding further management.

TBSRTC is a vital guideline which can bridge the communication gap and useful to maintain uniformity not on in the confined region but also worldwide. We followed the TBSRTC guidelines and every case was classified according to six tier reporting guideline from category I to VI.

Study includes total 160 cases of thyroid lesions which comprised of 140 cases (87.5%) of total. Published data suggest FNA has an overall accuracy rate around 75% in the detection of thyroid malignancy.^[9] The age group which was studied ranged from 5 years to 70 years and maximum no. of cases were in the age group 20-50 years means bulk of thyroid diseases were frequently encountered in young and middle aged group also the majority of cases were the females in reproductive age groups. In present study a female preponderance was noted. Similar female preponderance was noted by Unnikrishnan et al.^[10]

Neoplastic lesions were 11 cases (6.88%). The benign cyst consistent with thyroglossal cyst were 4 cases (2.06%) and others were 4 cases (2.06%). The bulk of the goitre cases were in the age group of 20-50 years and thyroiditis cases were in 11-40 years and mainly in the reproductive age group of the females.

Category I: This category includes cases in which sufficient material was not available like insufficient follicular cells (Satisfactory for evaluation: six groups of well visualised follicular cells with at least ten cells per group), cyst fluid only, obscuring blood, only macrophages, preparation artefact. In such cases repeat FNA was carried out under ultrasound guidance.

Category II: Majority of lesions were benign mainly of colloid goitre 97 cases (69.29%) out of 140. In comparison to various studiesbenign category includes 60-70% ^[8] reason for that is we have goitre belt here. Similar findings were observed Unnikrishnan et al. ^[10]The chances of thyroiditis after reproductive age appeared minimal from this study.

Category III: It is reserved for specimens that contain cells (follicular, lymphoid, or other) with architectural and/or nuclear atypia that is not sufficient to be classified as suspicious for a follicular neoplasm, suspicious for malignancy, or malignant. Management in such lesions is repeat FNAC after an appropriate interval. To be noted that this category is of last resort & should not be used indiscriminately.^[8]

Category IV: The goal of this category is to identify all potential follicular carcinomas and refer them for a diagnostic lobectomy. Although these cytomorphologic features do not permit distinction from a follicular adenoma (FA), they are reportable as Follicular Neoplasm (FN) or suspicious of Follicular Neoplasm (SFN), leading to a definitive diagnostic procedure. usually lobectomy. ^[11, 12, 13] The term SFN is preferred by some laboratories over FN for this category because a significant proportion of cases (up to 35%) prove not to be neoplasms but rather hyperplastic proliferations of commonly follicular cells. most those of multinodulargoiter. [14-17] About 15% to 30% of cases called FN/SFN prove to be malignant. [11, 14, 16] The majority of FN/SFN cases turn out to be FAs or adenomatoid nodules of multinodulargoiter, both of which are more common and outnumbers the Follicular carcinoma.

Category V: FNAC can diagnose many of the thyroid cancers with fair accuracy, especially papillary thyroid

carcinoma (PTC) which can be diagnosed with certainty by FNA. But the nuclear and architectural changes of some PTCs are subtle and focal. This is particularly true of the follicular variant of PTC, which can be difficult to distinguish from a benign follicular nodule. ^[18] Other PTCs may be incompletely sampled and yield only a small number of abnormal cells. ^[19] If only 1 or 2 characteristic features of PTC are present or if they are only focal and not widespread throughout the follicular cell population, or if the sample is sparsely cellular, a malignant diagnosis cannot be made with certainty. Such cases occur with some regularity, and they are best classified as suspicious for malignancy," qualified as "suspicious for papillary carcinoma." Such cases suspicious for papillary carcinoma are resected by lobectomy or thyroidectomy. Most (60%-75%) prove to be papillary carcinomas, and the rest are usually FAs. ^{[11,} ^{14, 16]} The same general principle applies to other thyroid malignancies like medullary carcinoma and lymphoma, but these are less frequent than PTC. Such cases were considered after correlating the other findings like serum calcitonin and calcium levels and other relevant data in medullary carcinoma.

Category VI: The malignant category is used whenever the cytomorphologic features are indicative of malignancy. After confirming the malignancy the sub classification was done after summarizing all the results. Approximately 3% to 7% of thyroid FNAs have conclusive features of malignancy, and most are papillary carcinomas.^[11, 14] Malignant lesions are usually treated bythyroidectomy, with some exceptions (e.g., metastatic tumors, non-Hodgkin lymphomas, and undifferentiated carcinomas). According to studies the positive predictive value of a malignant FNA interpretation is 97% to 99%.

We here practiced mainly the non-aspiration technique in almost all cases for studying cytology and found that it is better than aspiration technique. Aspiration technique is associated with low cellularity and more blood as compared to non-aspiration method. We recommend the non-aspiration method for FNAC of thyroid lesions. Similar suggestions by different studies like Maurya et.al [20] also recommend the non-aspiration technique better for thyroid lesion evaluation by FNAC. The study found thatit is difficult to differentiate follicular/Hurthle cell adenoma from carcinoma on cytological assessment because cytology cannot evaluate the criteria of vascular or capsular invasion or of intrathyroid spread. But the papillary carcinoma, Anaplastic carcinoma and medullary carcinoma can be diagnosed by characteristic cytological features.

V. Conclusion

TBSRTC is a vital guide for accurate management of thyroid lesions. Classifying the lesions in six categories and following the guidelines given by

The Bethesda USA meetings solves all problems regarding the management of thyroid lesions and leaves no confusion. It plays a big role in establishing the uniform communications between the managing medical personnel. Marked cellularity of the smear is the problem inherent in thyroid FNAC. Increased cellularity of the smear and loss of cohesion may be present in hyperplastic/adenomatous goiter and follicular neoplasm which causes difficulty in differentiating them. This can be solved by using The Bethesda System of Reporting thyroid lesions. We experienced that The Bethesda Reporting Systemisbest for management of thyroid lesions as it gives uniform reporting system.

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Age in	Male	Female	l	II		IV	V	VI	Total
years									
1-10	1	1	0	2	0	0	0	0	2
11-20	2	5	0	6	0	0	0	1	7
21-30	4	43	2	44	0	1	0	0	47
31-40	11	41	4	45	0	0	1	2	52
41-50	4	27	1	25	0	3	0	2	31
51-60	1	8	0	8	0	1	0	0	9
61-70	1	11	2	10	0	0	0	0	12
Total	24	136	9	140	0	5	1	5	160
	(15%)	(85%)	(5.63%)	(87.5%)	(0%)	(3.12%)	(0.63%)	(3.12%)	

Table 1: Distribution of thyroid lesions according to age and sex after applying TBSRTC

I-Non diagnostic or unsatisfactory, II- Benign, III-Atypia of undetermined significance or Follicular lesions of undetermined significance, IV- Follicular neoplasm or suspicious of follicular neoplasm, V- Suspicious of malignancy, VI- Malignant lesion.

 Table 2 : Diagnostic categories, implied risk of malignancy & recommended clinical management accordingTBSRTC. (Modified by Ali and Cibas)^[8]

Diagnostic category	Cytological diagnosis	Risk of malignancy	Clinical management
1	Non-diagnostic or	1-4 %	Repeat FNA with
	Unsatisfactory		ultrasound guidance
	Benign	0-3 %	Clinical follow-up
III	AUS/FLUS*	5-15 %	Repeat FNA
IV	FNS/SFN ¶	15-30 %	Surgical lobectomy
V	Suspicious of malignancy	60-75 %	Near total thyroidectomy
			or Surgical lobectomy
V	Malignant	97-99 %	Near total thyroidectomy

*AUS/FLUS – Atypia of undetermined significance / Follicular lesion of undetermined significance. ¶ FNS / SFN – Follicular neoplasm / Suspicious of follicular neoplasm.

Figure 1 : Abundant watery thin colloid in Colloid Goitre



Figure 2 : Hurthle cell changes in Hashimoto thyroiditis



Figure 3 : Numerous macrophages only: Nondiagnostic category



Figure 4 : Medullary carcinoma of thyroid with amyloid like material



Figure 5 : Follicular neoplasm with numerous microfollicles



Figure 6 : Anaplstic carcinoma of thyroid



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Determination of the Causative Agents of Bacteremia in Children under 5 Years and their Susceptibility Pattern to the Antibiotics

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Abstract- Objective: To determine the causative agents of bacteremia in children under 5 years and their susceptibility to the commonly used antibiotics.

Methods: This was a cross-sectional study on children (aged from 1 day to 5 years) admitted to the paediatric ward. The patients included all newborn babies and children admitted with fever and suspected of having sepsis. All the included children were clinically diagnosed for septicemia following strict aseptic precautions and the blood sample was taken. Blood culture were done by standard method.

Results: The overall incidence of bacteremia was 23.1%. The incidence of bacteremia was higher among the chidren of age group 13-60 month (38.1%) than <1 month (23.4%) and 1-12 month (12.5%). The male (25.8%) children were affected than females (19.6%).

Keywords: bacteremia, incidence, children, bacterial isolates.

GJMR-C Classification : NLMC Code: WC 240

DETERMINATION OF THE CAUSATIVE AGENTS OF BACTEREMIA INCHILDRENUNDER SVEARSANDTHEIR SUSCEPTIBILITY PATTERNTOTHEANT BIDTICS

Strictly as per the compliance and regulations of:



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Results: The overall incidence of bacteremia was 23.1%. The incidence of bacteremia was higher among the chidren of age group 13-60 month (38.1%) than <1 month (23.4%) and 1-12 month (12.5%). The male (25.8%) children were affected than females (19.6%). The *Klebsiella species* organism was the most common organism which was 44.4% followed by *Staphylococcus. Aureus* (14.8%) and *Coagulase Negative staphylococci (CONS)* (11.1%). The percentage of other organism was less than 10%. The percentage of gram negative bacterial isolate was among 70.4% of the samples and gram negative was 29.6%.

Conclusion: The resistance of the recovered *Klebsiella* spp. isolates to a number of antimicrobial agents was determined, and a pattern of multi-resistance was observed which may explain the prevalence of these isolates in pediatric bacteremia.

Keywords: bacteremia, incidence, children, bacterial isolates.

I. INTRODUCTION

acteremia is the presence of viable bacteria in the circulating blood. Bacteria may enter the blood stream giving rise to bacteremia from an existing focus of infection from a site with the commensally flora or by direct inoculation of contaminated materials into the vascular system. These organisms are often cleared from the blood within minutes, so the bacteremia is silent and transient, but if the immune system is overwhelmed or evaded, organisms persist in the blood and bacterimic symptoms would arise. Bacteremia should be distinguished from septicemia in which signs and symptoms of severe diseases are present¹. Neonates are particularly vulnerable to infections

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because of their weak immune barrier. Several risk factors have been identified both in the neonates and children which makes them susceptible to infections². Children with septicaemia present with fever, difficulty in breathing, tachycardia, malaise, refusal of feeds or lethargy³.

Studies of bloodstream infections in children admitted to African hospitals suggest that the prevalence of bacterial bloodstream infections among inpatients with fever or clinical sepsis exceeds that described in wealthier regions^{4,5}. Bloodstream infections continue to be the major cause of morbidity and mortality despite advance in antimicrobial therapy and supportive care⁶. Fever in infants younger than 1 year old, especially those younger than 3 months, can signal a serious infection⁷.

The aim of this study was to determine the causative agents of bacteremia in children under 5 years and their susceptibility to the commonly used antibiotics.

II. MATERIAL AND METHODS

This was a cross-sectional study approved by the ethical committee of the institute. In this study, 117 blood samples were collected from children (aged from 1 day to 5 years) admitted to the paediatric ward of a teaching hospital in north India. The patients included all newborn babies and children admitted with fever and suspected of having sepsis. Children with fever less than 5 days and with known clinical condition such as malignancies, tuberculosis etc. were excluded from the study. The consent was taken from parent/guardian of each children before enrolling in the study.

A total of 128 children were included in this study. All the included children were clinically diagnosed for septicemia following strict aseptic precautions and the blood sample was taken. One milliliter (neonates) and 5 ml (children) blood were collected, inoculated into 10 ml and 50 ml, respectively of brain heart infusion broth. The culture bottles were incubated at 37°C aerobically and periodic subcultures were done onto Mac Conkey's agar, blood agar and chocolate agar after overnight incubation on day three, day four and finally on day growth obtained was seven⁸. The identified by conventional biochemical tests and the antibiotic sensitivity testing was performed on Mueller-Hinton agar plates by Kirby-Bauer disc diffusion method. Zone

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diameter was measured and interpreted as per the Clinical and Laboratory Standards Institute (CLSI) guidelines as used by Tiwari et al⁸. Bacterial sensitivity was tested for the following antimicrobials: amikacin, amoxicillin clavulanic acid, ampicillin, aztreonam, cefotaxime, ceftazidime, ceftriaxone, cephalexin, cefoxitin, ciprofloxacin, gentamicin, imipenem, meropenem, piperacillin-tazobactam, tobramycin, linezolid and vancomycin.

Methicillin resistance in Staphylococcus aureus (MRSA) was tested using Mueller-Hinton agar with 4% NaCl with cefoxitin disc (30 micrograms) by Kirby-Bauer disc diffusion method. A zone size of >22 mm was considered sensitive and < 21 was considered as Suspected extended-spectrum resistant. beta lactamases (ESBLs) producing organisms were confirmed by double disk synergy test⁸. Detection of plasmid-mediated AmpC was done by the AmpC disk test and the isolates showing reduced susceptibility to carbapenems (imipenem and meropenem) were selected for detection of metallo-beta lactamases (MBLs) enzymes by imipenem-EDTA disk method. For quality control of disc diffusion tests ATCC control strains of E. coli ATCC 25922, S. aureus ATCC 25923 and P. aeruginosa ATCC 27853 strains were used. The results were expressed as percentages. Microsoft excel was used for the interpretation of these results.

III. Results

Table-1 presents the percentage of children of Bacteremia according to demographic profile. The overall incidence of bacteremia was 23.1%. The incidence of bacteremia was higher among the chidren of age group 13-60 month (38.1%) than <1 month (23.4%) and 1-12 month (12.5%). The male (25.8%) children were affected than females (19.6%).

The Klebsiella species organism was the most common organism which was 44.4% followed by *Staphylococcus. Aureus* (14.8%) and *Coagulase Negative staphylococci (CONS)* (11.1%). The percentage of other organism was less than 10% (Fig.1).

The percentage of gram negative bacterial isolate was among 70.4% of the samples and gram negative was 29.6% (Fig.2).

Table-2 depicts antibacterial resistance pattern of the gram negative blood stream isolates. Klebsiella spp organism was resistance to ampicillin (3), amoxyclav (5), amikacin & gentamycin (2) and one each tocotrimoxazole, ciprofloxacin, cefotaxime& aztreonam. The frequency of other organism resistance to most of drugs was one.The frequency gram positive isolates resistance to most of the drugs was one (Table-3).

IV. Discussion

Who guidelines for management of acute illness in children (Integrated Management of Childhood

Illness) recommend use of an appropriate antibacterial drug in addition to an antimalarial drug in children with certain signs of severe illness⁹. Despite considerable progress in hygiene, antimicrobial therapy, and supportive treatment, blood stream infections remain important causes of morbidity and mortality which, may reaches to 20%-30%¹⁰. Microbiologic culturing of blood is the only available means for diagnosis of these infections and allows for successful recovery of bacteria in 99% in patients with bacteremia of septicemia¹¹. An American review covering a 50-years period has show major changes in the etiology of neonatal septicemia¹².

In this study, the incidence of bacteremia was higher among male children. Similar findings had been reported in southern state of India⁸. Some other studies had also reported higher incidence of bacteremia in male children^{13, 2}. The incidence of bacteremia was higher among 13-60 month (38.1%) in this study which was contradictory to the studies by Tsering et al¹⁴ and Meremkwer et al.² in which incidence of bacteremia was more common among newborns. In this study, the culture positivity rate was found to be 23.1% (23/117). Almost similar rate had been reported in a Indian study in which blood culture positivity rate of 25%⁸. Other studies had also reported similar positivity rate^{15, 14}.

The percentage of gram negative organism found in this study was similar to other studies also^{16,15}. In the present study, the *Klebsiella* species was the commonest isolate associated with bacteremia which was similar to study by Al-Charrakh et al¹⁷. Many studies have been shown that Gram positive organisms were the mainly *Staphylococcus aureus* as the most frequently isolated bacteria causing bacteremia^{18,19}.

There was varying number of the gram negative and positive organisms resistance to different drugs in this study. Prabhu et al²⁰ reported that the gram negative organisms showed maximum resistance to ampicillin. However, Tiwari et al⁸ reported that the gram positive organisms showed 77.78% resistance to penicillin but were 100% sensitive to linezolid and vancomycin. Among the 6 *Staphylococcus aureus*, 2(33.33%) were detected as Methicillin resistant *Staphylococcus aureus* (MRSA).

One of the limitation of this study is lesser sample size. Studies on larger sample size is recommended for better interpretation of the results. The studies on the community acquired blood stream infections is also needed at present.

V. Conclusion

The resistance of the recovered *Klebsiella* spp. isolates to a number of antimicrobial agents was determined, and a pattern of multiresistance was observed which may explain the prevalence of these isolates in pediatric bacteremia.

Conflict of interest: None

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	No. of	Bacteremia			
	children	No. %			
	assessed				
Age group					
<1 month	64	15	23.4		
1-12	32	4			
months			12.5		
13-60	21	8			
month			38.1		
Gender					
Male	66	17	25.8		
Female	51	10	19.6		
Total	117	27	23.1		

Table 1 : Percentage of children of Bacteremia according to demographic profile

Table	2 : Antibacterial	resistance pat	tern of the	aram neaative	blood stream isola	ates

Antibiotics	Klebsiella spp (n=12)		E.coli	xoli (n=2) Pse aer (n=		Pseudomonas aeruginosa (n=2)		tobacter anii (n=1)	S.typł (n=2)	ni '	Citrob freund (n=1)	bacter dii
	No.	%	No.	%	No.	%	No.	%	No.	%	No.	%
Ampicillin	3	25.0	1	50.0	NT	-	NT	-	2	100. 0	1	100.0
Amoxyclav	5	41.7	0	0.0	NT	-	NT	-	1	50.0	0	0.0

Amikacin	2	16.7	1	50.0	1	50.0	0	0.0	0	0.0	1	100.0
Cotrimoxaz	1	8.3	1	50.0	NT	-	0	0.0	1	50.0	0	0.0
ole												
Gentamycin	2	16.7	0	0.0	1	50.0	1	100.0	0	0.0	0	0.0
Tobramycin	NT	-	NT	-	1	50.0	0	0.0	NT	-	NT	-
Ciprofloxaci	1	8.3	0	0.0	2	100.0	1	100.0	1	50.0	1	100.0
n												
Cefotaxime	1	8.3	1	50.0	NT	-	NT	-	1	50.0	0	0.0
Ceftriaxone	2	16.7	0	0.0	NT	-	NT	-	0	0.0	0	0.0
Ceftazidime	NT	-	NT	-	0	0.0	1	100.0	NT	-	NT	-
Aztreonam	1	8.3	0	0.0	0	0.0	0	0.0	NT	-	0	0.0
Imipenem	0	0.0	0	0.0	0	0.0	0	0.0	0	0.0	0	0.0

NT-Not Tested

Table 3 : Antibacterial resistance pattern of gram positive blood stream isolates

Antibiotics	Staphylococcus aureus (n=4)	Staphylococcus aureus (n=4)		
	No.	%	No.	%
Penicillin	2	50.0	1	33.3
Amoxyclav	1	25.0	1	33.3
Cefoxitin	1	25.0	2	66.7
Erythromycin	2	50.0	1	33.3
Linezolid	1	25.0	0	0.0
Vancomycin	0	0.0	1	33.3



Figure 1 : Distribution of organisms isolated from blood culture



Figure 2 : Prevalence of gram negative and gram positive bacterial isolates

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A Study on Cold Agglutinins in Malaria from a Tertiary Care Hospital of South India

By Venkatesh V. N, Ravish Kumar M & Nagendra Gowda M. R

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Abstract- Background: Malarial infection associated with common hematological abnormalities like anemia, Thrombocytopenia, Hemolytic anemia. Hemolytic anemia is due to occurrence of cold agglutinins. This study throws light on the prevalence of cold agglutinins in patients with malarial infection and their detection.

Method: About 150 patients diagnosed as having Malaria by Peripheral smear, QBC for malaria or by Rappid diagnostic test for Malaria were studied for occurrence of cold agglutinins by Cold agglutinin test.

Results: Out of 150 Patients, who are diagnosed as positive for malaria, 83.3% of the patients had negative for cold agglutinin and only (4%) were found to have high titers of cold agglutinins. The mean age group of the patients who were positive for the test was 45.25 ± 19.7 years. The Hemoglobin percentage and platelet count in cold agglutinin positive patients were 8.85 (\pm 3.72) gms% and 1, 03, 500 (\pm 1.16) respectively.

GJMR-C Classification : NLMC Code: WC 750, QV 256

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A Study on Cold Agglutinins in Malaria from a Tertiary Care Hospital of South India

Venkatesh V. N °, Ravish Kumar M ° & Nagendra Gowda M. R^P

Abstract- Background: Malarial infection associated with common hematological abnormalities like anemia, Thrombocytopenia, Hemolytic anemia. Hemolytic anemia is due to occurrence of cold agglutinins. This study throws light on the prevalence of cold agglutinins in patients with malarial infection and their detection.

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Conclusion: The prevalence of cold agglutinins in patients infected with Plasmodium vivax or P falciparum was 2.67%. There was a significant change in hematological parameters like hemoglobin but not platelets.

I. INTRODUCTION

alaria is a public health problem in Sub Saharan Africa, some parts of South East Asia and South America with considerable morbidity and mortality. This disease was almost eradicated in India during 1960's but to reemerge as a public health problem in last few decades¹. This is mainly due to development of resistance by mosquitoes against the insecticides and to therapeutic agents by the plasmodium organism. The social and environmental changes also plays major role in this situation. Usually the disease malaria results from the parasites belonging to Plasmodium species. Human malaria is a result of four different species of plasmodium including Vivax, Falciparum, Ovale and Malariae. Nowadavs. Plasmodium Knowlesii a fifth parasite which was known cause malaria in monkeys has been implicated in some parts of Karnataka.

Simple techniques including examination of the peripheral smear is used to establish the diagnosis in Malaria. Fluorescent technique can also be employed to detect the malarial parasite. Nowadays, Malaria is often considered as grey area between the parasitology and hematology. Recent text books have considered malaria as a typical example of Hemolytic Anemia as result of acquired extra corpuscular causes. The common hematological abnormalities accompanied with the malarial infection are anemia, thrombocytopenia, spleenomegaly, mild to moderate atypical lymphocytosis and rarely DIC.²

The available literature had shown that, the main reason for hemolytic anemia especially in malaria is due to occurrence of cold agglutinins. The cold agglutinins are capable of agglutinating RBCs and hence result in hemolysis. But the literature available shows that the cold agglutinins occur transiently in malaria. This is mainly due to activation of polyclonal B lymphocytes which is predominantly of IgM variety. Rarely might they be either IgA or IgG variety which has the specificity for blood group antigen. The literature suggests that the hemolysis appears in 2 - 3 weeks after the malarial infection. The hemolysis is usually mild and self limiting and occasionally severe and fatal.^{3, 4}

Increasing emphases is now given for these antibodies in protection against human malaria which is directed at erythrocytic stages of Plasmodium falciparum. But the protection offered by such antibodies is relatively unstable and the precise role of specificities remains unclear regarding the antigenic variability of parasite proteins. Even though parasite specific antibodies formed contribute to protection, it is not evident to what extent antibodies so formed contribute to protection⁵. Many studies have shown that these antibodies result in hemolytic anemia due to complement mediated RBC destruction in the reticuloendothelial system.

Studies regarding role of cold agglutinins are scant in India and World. This made us to take up this study in order study the profile of patients with Cold agglutinins in Malaria.

II. MATERIALS AND METHODS

This cross sectional study was undertaken in the Department of Microbiology, Basaveshwara Medical College and Hospital, Chitradurga. About 150 patients who were diagnosed as having Malaria by peripheral

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smear examination (both thick and thin), MPQBC or by malarial antigen assay before starting the antimalarial treatment were included in the study. An informed consent was obtained from each patient before the study was started and clearance from Intuitional ethical committee was obtained. This study was carried out between January, 2012 to March, 2013.

A detailed history was taken followed by detailed clinical examination for all the patients included in the study. These patients were also investigated for Hemoglobin estimation by cyanmethemoglobin method and Total platelet count by modified Dacie Leurs method.

Cold agglutinin test was used to detect cold agglutinins in the malarial patients. The cold agglutinins if present in patient's serum in high titers may be pathologic and result in cold agglutinin disease. Serum or plasma of all the malarial patients was separated at 37° C from a Blood sample collected. A pool of 2 or

more examples of washed group O1 adult red cells and Phosphate buffered saline (PBS) at pH 7.3 were used as reagents. A serial two fold dilutions of the patient's serum or plasma in PBS were prepared followed by Two drops of each dilution 1 drop of a 3% to 5% suspension of red cells was mixed. The solution was mixed and incubated at 4° C for 1 to 2 hours.

The tubes were centrifuged for 15 to 20 seconds at 900 to 1000 X g. Then it was placed in ice water bath. The tubes were examined microscopically for agglutination, starting with the tube at highest dilution. The results were graded and recorded.

All the details were collected in a pre-structured, self administered proforma. The data thus obtained was compiled and analysed in the form of frequency and proportions. Chi square test and Student T test were used as significance test by using Statistical Package for Social Sciences (SPSS vs 18). A p value of less than 0.05 was considered as statistically significant.

III. Results

Table 1 : Cold agglutinin test results of the study group

	Frequency	Percent
Negative	125	83.3
Positive at 4 deg at 1/256 dilution	3	2.0
Positive at 4 deg at 1/64 dilution	1	0.7
Positive at 4 deg at 1/32 dilution	3	2.0
Positive at 4 deg C at 1/16 dilution	5	3.3
Positive at 4 deg at 1/8 dilution	10	6.7
Positive at 4 deg at 1/4 dilution	3	2.0
Total	150	100.0

On observing the results cold agglutinin test, about 83.3% of the patients had negative cold agglutinin test. Only 4 patients who had malaria were positive for the cold agglutinin test since the cut of value for the positive test was above 1:64 dilution.

Table 2 : Distribution of the Malarial patients according to age group and result of Cold agglutinin test

Age group	Cold agglutinin t	Cold agglutinin test		
	Negative n (%)	Positive n (%)	n (%)	
Less than 20 yrs	25 (17.1)	1 (25.0)	26 (17.3)	
21 - 30 yrs	57 (39.0)	0	57 (38.0)	
31 - 40 yrs	25 (17.1)	0	25 (16.7)	
41 - 50 yrs	17 (11.6)	2 (50.0)	19 (12.7)	
51 - 60 yrs	10 (6.8)	0	10 (6.7)	
61 yrs and above	12 (8.2)	1 (25.0)	13 (8.7)	
Total	146 (100)	4 (100)	150 (100)	
Mean ± Std dev	34.09 ± 15.55	45.25 ± 19.7	34.39 ± 15.7	

t (148)= 1.406 p=0.162, NS

Table no 2 shows the age group wise distribution of the study group. The mean (\pm SD) age of subjects in the study group was 34.39 \pm 15.7 years. The mean age group of the patients who were positive for the test was 45.25 \pm 19.7 years. About 50% of the

subjects who had positive test belonged to 51 - 60 years age group, one patient was aged less than 20 years and another aged more than 60 years. There was no statistically significant difference between the age of the patients and result of the Cold agglutinin test.

Table 3 : Distribution of the Malarial patients according to sex and result of Cold agglutinin test.

	Cold aggluti	Total		
Sex	Negative n (%)	Positive n (%)	n (%)	
Male	104 (71.2)	3 (75.0)	107 (71.3)	
Female	42 (28.8)	1 (25.0)	43 (28.7)	
Total	146 (100)	4 (100)	150 (100)	
χ^2 value = 0.027 df=1 p=0.869				

About 75.0% of the patients who were positive for the cold agglutinin test were males and 25% were females. Of 146 patients who had shown negative results for the test, 71.2% were males and 28.8% were females. This difference was not statistically significant.

Table 4 : Distribution of the Malarial patients according to the level of hemoglobin and result of Cold agglutinin test.

Haemoglobin %	Cold agglutinin	Total	
	Negative n (%)	Positive n (%)	
More than normal	35 (24.0)	2 (50.0)	37 (24.7)
Less than normal	111 (76.0)	2 (50.0)	113 (75.3)
Total	146 (100)	4 (100)	150 (100)
Mean ± Std dev	11.40 ± 2.67	8.85 ± 3.72	11.33 ± 2.72

t (148)= 1.862

p=0.065, NS

The mean hemoglobin level of the study subjects was 11.33 (\pm 2.72) gm%. Patients negative cold agglutinin test results had a mean hemoglobin level of 11.4 (\pm 2.67) gm% and those with positive test had a mean hemoglobin level of 8.85 (\pm 3.72) gm%. About

50% of the patients had normal level of hemoglobin and 50% had less than normal level of hemoglobin. There was statistically significant difference in the hemoglobin levels between the negative and positive cold agglutinin test patients.

Table 5 : Distribution of the malarial patients according to the level of Platelets and result of Cold agglutinin test

Platelet Count	Cold agglutinin test	Total	
	Negative n (%)	Positive n (%)	11 (%)
Less than 1,65,000	132 (90.4)	3 (75.0)	135 (90.0)
Normal	13 (8.9)	1 (25.0)	14 (9.3)
More than 4,15,000	1 (0.7)	0	1 (0.7)
Total	146 (100)	4 (100)	150 (100)
Mean \pm Std dev	97,832.2 ± 63,824	1,03,000 ± 1.16	97,970.0 ± 65,099.4

t (148)= 0.156 p= 0.876, NS

The mean platelet count in test negative patients was 97,832.2 (\pm 63,824) and in test positive patients was 1, 03,500 (\pm 1.16). This difference was not

statistically significant. The platelet count was low in 91.4% of test negative and 75% of the test positive patients.

Table 6: Distribution of the Malarial patients according to the type of malarial parasite and result of Cold agglutinin test

Type of Malarial Parasite	Cold agglutir	Total n (%)	
	Negative n (%)	Positive n (%)	
Pl. Vivax	98 (67.1)	0	98 (65.3)
PI. Falciparum	32 (21.9)	2 (50.0)	34 (22.7)
Mixed	16 (11.0)	2 (50.0)	18 (12.0)
Total	146 (100)	4 (100)	150 (100)
$\chi^2 = 8.985$	df=2	p=0.01	1, Sig

Table no 6 shows the type of malarial parasitic infection and results of the Cold agglutinin test. About 67.1% of the test negative patients were infected with Plasmodium Vivax. About 50% of the malarial patients with positive cold agglutinin test were infected with Plasmodium vivax and 50% had mixed Vivax and Falciparum infection. The difference in type of parasite and results of the Cold agglutinin test was statistically significant.

IV. Discussion

This study was mainly undertaken to study the prevalence of cold agglutinins in malaria patients. The results from this study had shown that about 2.67% of the malarial patients were positive for cold agglutinins with titres above 1:64 dilutions. In a study in 1980 the authors have failed to demonstrate any cold agglutinins in 1980.⁶ A study of monoclonal gammopathies of cold agglutinin disease in mayo clinic had shown that the prevalence was 1.1%. A study by Torres et al had shown prevalence similar to that observed in this study.⁷

The mean age group of the patients who were positive for the test was 45.2 (\pm 19.7) years. About 50% of the patients who were positive for the test belonged to 41 – 50 years and 61 years and above age group. However there was a statistically significant difference between age group and result of the Cold agglutinin test. The finding from our study shows that the prevalence of cold agglutinins was common in the patients above 40 years. In contrary to these findings, Sharon⁴ states that the secondary cold agglutinin disease is common in children and young adults. About 75% of the patients who had cold agglutinins were males in this study.

The mean haemoglobin levels in the positive patients for cold agglutinins was 8.85 (\pm 3.72) gm% in this study. About 50% of the patients had hemoglobin more than normal and 50% had hemoglobin below normal. This study is able to demonstrate haemolysis similar to the study by Gertz et al¹⁸. There was no drop in platelet count in patients who were positive for cold agglutinins in this study. Reduced levels of Haemo-

globin and Platelet count in malaria patients were recorded by other author ⁸.

About 67.1% of the test negative patients were infected with Plasmodium Vivax. About 50% of the patients' positive for Cold agglutinin test had mixed malarial parasitic infections. The difference in type of parasite and results of the Cold agglutinin test was significant. A study from Ouagadougou also reported the similar results.

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33. Report concluded results: Use concluded results. From raw data, filter the results and then conclude your studies based on measurements and observations taken. Significant figures and appropriate number of decimal places should be used. Parenthetical remarks are prohibitive. Proofread carefully at final stage. In the end give outline to your arguments. Spot out perspectives of further study of this subject. Justify your conclusion by at the bottom of them with sufficient justifications and examples.

34. After conclusion: Once you have concluded your research, the next most important step is to present your findings. Presentation is extremely important as it is the definite medium though which your research is going to be in print to the rest of the crowd. Care should be taken to categorize your thoughts well and present them in a logical and neat manner. A good quality research paper format is essential because it serves to highlight your research paper and bring to light all necessary aspects in your research.

INFORMAL GUIDELINES OF RESEARCH PAPER WRITING

Key points to remember:

- Submit all work in its final form.
- Write your paper in the form, which is presented in the guidelines using the template.
- Please note the criterion for grading the final paper by peer-reviewers.

Final Points:

A purpose of organizing a research paper is to let people to interpret your effort selectively. The journal requires the following sections, submitted in the order listed, each section to start on a new page.

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- Separating a table/chart or figure impound each figure/table to a single page
- Submitting a manuscript with pages out of sequence

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The summary should be two hundred words or less. It should briefly and clearly explain the key findings reported in the manuscript-must have precise statistics. It should not have abnormal acronyms or abbreviations. It should be logical in itself. Shun citing references at this point.

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- Fundamental goal
- To the point depiction of the research
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- Significant conclusions or questions that track from the research(es)

Approach:

- Single section, and succinct
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Approach:

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- If use of a definite type of tools.
- Materials may be reported in a part section or else they may be recognized along with your measures.

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Approach:

- It is embarrassed or not possible to use vigorous voice when documenting methods with no using first person, which would focus the reviewer's interest on the researcher rather than the job. As a result when script up the methods most authors use third person passive voice.
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What to keep away from

- Resources and methods are not a set of information.
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The principle of a results segment is to present and demonstrate your conclusion. Create this part a entirely objective details of the outcome, and save all understanding for the discussion.

The page length of this segment is set by the sum and types of data to be reported. Carry on to be to the point, by means of statistics and tables, if suitable, to present consequences most efficiently. You must obviously differentiate material that would usually be incorporated in a study editorial from any unprocessed data or additional appendix matter that would not be available. In fact, such matter should not be submitted at all except requested by the instructor.



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- Sum up your conclusion in text and demonstrate them, if suitable, with figures and tables.
- In manuscript, explain each of your consequences, point the reader to remarks that are most appropriate.
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• Examine your data, then prepare the analyzed (transformed) data in the form of a figure (graph), table, or in manuscript form. What to stay away from

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Approach

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- Give details all of your remarks as much as possible, focus on mechanisms.
- Make a decision if the tentative design sufficiently addressed the theory, and whether or not it was correctly restricted.
- Try to present substitute explanations if sensible alternatives be present.
- One research will not counter an overall question, so maintain the large picture in mind, where do you go next? The best studies unlock new avenues of study. What questions remain?
- Recommendations for detailed papers will offer supplementary suggestions.

Approach:

- When you refer to information, differentiate data generated by your own studies from available information
- Submit to work done by specific persons (including you) in past tense.
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Methods and Procedures	Clear and to the point with well arranged paragraph, precision and accuracy of facts and figures, well organized subheads	Difficult to comprehend with embarrassed text, too much explanation but completed	Incorrect and unorganized structure with hazy meaning
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Discussion	Well organized, meaningful specification, sound conclusion, logical and concise explanation, highly structured paragraph reference cited	Wordy, unclear conclusion, spurious	Conclusion is not cited, unorganized, difficult to comprehend
References	Complete and correct format, well organized	Beside the point, Incomplete	Wrong format and structuring

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