

Exploring biofilm forming capacity of *A.baumannii* from different clinical sources and screening copper nanoparticles for antibiofilm activity.

By

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Project report completed under the supervision of

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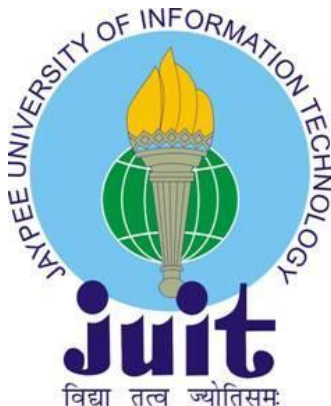
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Jaypee University of Information Technology



CERTIFICATE OF ORIGINALITY

This is to certify that the work titled " Exploring biofilm forming capacity of *A.baumannii* from different clinical sources and screening copper nanoparticles for antibiofilm activity" submitted by Chetansee Khanna in fractional completion of the necessities for the award of degree of Bachelors of Technology in Biotechnology, of Jaypee University of InformationTechnology, Solan, has been approved underneath the direction of Dr. Jitendraa Vashistt. This work has not been presented partially or completely to any other University or Institute for the grant of this or any other degree or diploma.

Signature of Supervisor

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Signature and name of the student

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Summary

A total of 48 *A. baumannii* strains were collected from Department of Microbiology (Bacteriology division), AIIMS, New Delhi during the study period. Clinical source of isolation of *A. baumannii* included patient's pus, respiratory and peritoneal fluid etc. Collected strains were characterized by standard biochemical assays (carbohydrate and amino acid utilization). *A. baumannii* (ATCC 19606) which is a type strain was also made in use as control against antimicrobial susceptibility analysis and biofilm production assay. Antimicrobial susceptibility test (AST) and minimum inhibitory concentration (MIC) of different antibiotics against isolates of *A. baumannii* were performed according to CLSI guidelines 2015 and ICMR standard operating procedure, bacteriology 2015. AST and MIC were done using Kirby Bauer's disk diffusion method and E- strips test, respectively.

Motility assays were performed which includes swarming and twitching to observe that if the pathogen is motile or not and to what extent motility is helping in dispersal of the pathogen in new environment. Antibiofilm activity of copper nanoparticles synthesized by green synthesis, amalgam with catharanthus-rosesus etc. was screened by antibiofilm assay.

Chapter 1: Introduction

Acinetobacter baumannii is a pleomorphic, aerobic, gram-negative coccobacillus bacterium. A health care related pathogen, which is also an opportunistic pathogen in human being, specially effecting people with immune-compromised systems due to any medical reasons [1]. These bacteria are normally correlated with water body environments [2], also it has been revealed to live and colonize on the skin as well as are now separated in huge amounts from the respiratory and oropharynx secretions of infected persons [3]. Now a days it is also exemplified as Multidrug-resistant and antimicrobial-resistant which is becoming the major reason for nosocomial (hospital derived) infections .Whereas to the scientists all over the world, its natural habitat is still indefinite [4]. From few years, it is renamed as “red alert” human pathogen, producing alertness among the medical alliance, occurring mainly due to wide and broad resistance spectrum of antibiotics [5]. With advancement in time *Acinetobacter baumannii* has appeared as a noteworthy hospital derived nosocomial pathogen, which is showing resistance to usually prescribed antimicrobials. It has newly expanded unsavory reputation as it causes some types of soft tissue infections in army men fighting and returning back from Iraq and Afghanistan. This occurrence of multidrug-resistant (MDR) pathogens has gradually emerged as solely the reason for various nosocomial and community-acquired infections [6]. The most common target of this organism is usually the pre-admitted patients in already immune-compromised state. Pneumonia which is derived from hospitals holds the most number of cases being reported due to this organism.

MDR characteristic is shown against many of the subsequent drug categories: antipseudomonal carbapenems (imipenem or meropenem), antipseudomonal cephalosporins (ceftazidime or cefepime), aminoglycosides (gentamicin, tobramycin, or amikacin), ampicillin-sulbactam and fluoroquinolones (ciprofloxacin or levofloxacin).

A Dutch microbiologist, named Martinus Willem Beijerinck, in 1911 revealed an aerobic, gram-negative, non-fermentative bacterium which we are acquainted with, is included in the genus *Acinetobacter* [7]. *Acinetobacter* started to be reported as a major nosocomial pathogen in the 1970s, though during those periods it was effortlessly treated due to its susceptibility towards prescribed antimicrobials. Whereas in the 1970s *A.baumannii* was listed to be sensitive to the majority antimicrobials, nowadays this organism displays broad spectrum resistance mechanism

to all most every first-line antibiotics [8]. Freshly, due to its apprehension in war regions, it has earned itself a label as “Iraqibacter”.

Usually, it sources outbreak of infection and health issues, together with wound infection, meningitis, pneumonia, urinary tract infection, and bacteremia [9]. Typically bacteria demonstrates motility by presenting twitching and swarming movements at times, other optional explanations for its motility is oozing of exopolysaccharide, which generates a film of elevated molecular weight sugar chains behind the bacterium to execute locomotion [10].

Chapter 2: Review of literature

Innate habitat:

Associates of the genus *Acinetobacter* are measured to be omnipresent organisms. Former verdicts have throw in to the frequent delusion that *A. baumannii* is also ever-present in nature [11]. The reality is, each kind of the genus *Acinetobacter* is not located in local environment but also majority of human extracted samples of *Acinetobacter* species always have function as pathogens [12, 13].

Epidemiology:-

Australia along with pacific islands

Primary reports of *A. baumannii* of this region showed community gained diseases, which have greatly diverse occurrence as compared to hospital-acquired illnesses. Indigenous racial setting, alcoholism, chronic obstructive airway disease, cigarette smoking and diabetes mellitus are being significant jeopardy issues. In addition to this, these mentioned strains are considerably vulnerable to antibiotics. Throat infections as well as micro-aspiration are implicated for pathogenicity of *Acinetobacter* caused infections [14, 15].

North America

There exist past cases of MDR reported happenings in this region. The organisms in this outburst were multidrug resistant, keeping hold of vulnerability only to polymyxins and ampicillin-sulbactam [16]. National supervision studies have established major method for development of MDR *Acinetobacter* species [17].

Africa

Statistics of resistance of this pathogen to various antimicrobials are usually for South Africa only, even though there are few reports from various countries [18]. Strains showing resistance are commonly found in few places (for an instance, burned regions, ICU wards etc.) which are being spread from one institution to other [19].

Latin America

Frequencies of non-susceptibility to various antibiotics like imipenem, gentamicin, ceftazidime, ciprofloxacin, meropenem and piperacillin-tazobactam in Latin America emerge to be amongst the uppermost issues [20]. A diversity of carbapenemases has also been recognized in *A. baumannii* isolates [21, 22, and 23].

Europe

Infections due *A.baumannii* have been a matter of concern in this country [24]. From 1980s, molecular typing methods specific to some regions are being used for detecting the hospital outbreaks of this pathogens infections [25]. Spreading of these strains is usually reported due to transfer of pre-infected patients from one hospital to another [26, 27].

Asia with the Middle East

Several eruption of pandrug-resistant species of *A.baumannii* are acknowledged from Asian along with some hospitals of Middle Eastern regions[28], also a range of carbapenemases are being depicted to be initiated from this region [29]. Resistance to tigecycline [30] along with polymyxin B [31] previously exists in this section.

Clinical manifestations of Acinetobacter baumannii infections

Pneumonia attained from hospital settings:

The majority of *A. baumannii* cases and the isolates derived from them are usually extracted from the respiratory routes of hospital admitted people. In most of the times, true pneumonia is difficult to be differentiated and characterized from other higher airway colonization infections. It also has been proved that ventilator-related pneumonia (VRP) is caused by this pathogen only. Classically, patients who have resided for long periods in ICU environment

are infected with *A. baumannii* infections [32], but it is contradictory in epidemic situations and conditions.

Community-gained Pneumonia

Many sultry sections of Australia as well as Asia have reported many cases of this type of pneumonia [33]. The disease most characteristically happens throughout the monsoon amid population with usual habit of alcoholism maltreatment which might cause necessary admittance in hospitals. This type of pneumonia is characterized by a lesser frequent route, lesser bloodstream related infections, and death rate ranging from 40 to 60% [34]. General cause of this infection would be throat infections, which happens in local population of around 10% with disproportionate alcoholism habits [35].

Meningitis

This type of disease is of nosocomial origin, which is also called as post-neurosurgical *A. baumannii* meningitis. More gram-negative pathogens are related in causing this disease in major of the cases [36], which makes it obvious for multidrug-resistant *A. baumannii* strains to be included in the list [37].

Traumatic Battlefield and Other Wounds

A. baumannii possibly, sporadically source either skin or soft tissue infections and diseases exterior of the military populace. These creatures caused 2.1% of ICU-acquired skin or soft tissue ailments in single evaluation [38]. It is a renowned pathogen in blaze units and could be tricky to eliminate from such patients [39]. *A. baumannii* is frequently cut off from wounds of war cases from Iraq as well as Afghanistan [40]. It was the mainly frequently secluded organism (32.5% of cases) in one evaluation of fight fatalities with usual and many tibia fractures.

Bloodstream Infection

A. baumannii was an additional reason of occurrence of ICU-derived blood infections as compared to non-ICU derived infection. This above mentioned *A.baumannii* bloodstream infection had the third peak rough mortality pace in the ICU. These types of infections were

the most recent of the entire cases of infections to happen throughout the medical environment [41]. Consequently it is not convinced whether the elevated death speed symbolizes its incidence in patients with continuing grave sickness or maybe this organism posses motility features which are not known.

Urinary Tract Infection

A. baumannii is an infrequent reason of UTI, which is accountable for just few cases of ICU-possessed UTIs [42]. Characteristically, *A.baumannii* is related with catheter-associated infection which is also commonly known as colonization. It is not standard for this organism to originate straightforward UTI in healthy individuals.

Other Manifestations

A little figure of occurrence of *Acinetobacter* endocarditis has also been reported [43]. The majority events have engaged prosthetic surgeries. *Acinetobacters* might source endophthalmitis or keratitis, many a times, which is linked to contact lens utilization or subsequent to any sort of eye surgery [44].

Frequency

A.baumannii is also called as 'Iraqibacter' owing to its apparently sudden appearance in armed curative amenities throughout the Iraq combat. It has sustained to be a subject of worry for veterans and soldiers who doled out in Iraq as well as those in Afghanistan. Additional with transitory time, multidrug-resistant *A. baumannii* has extended to national hospitals in division owing to the transport of infected soldiers during diverse medical services [45]. From precedent years diverse surveys, it was affirmed that infection do not happen because of this bacteria at the instance of injury only, but is further probable nosocomially obtained, predictably because of the capacity of *A. baumannii* to continue on synthetic and abiotic facades for extensive stages, and the numerous conveniences to which wounded soldiers are uncovered throughout the casualty-clearance course and agenda [46, 47]. Risk reasons which adds to the possibilities of infection are protracted span of hospital wait, revelation to an intensive care unit (ICU), receiving of

perfunctory aeration, colonization strain, revelation to antimicrobial factors, existing operation, all-inclusive measures, and original severity of infection [47, 48].

Widespread of environmental infectivity is frequently illustrated, and outbursts of infection have been drawn to respiratory gears, injury treatment tasks, humidifiers, and patient concerned stuff [49, 50]. From diverse surveys the multi drug-resistance bacterial infections are normally established on a diversity of medical gears, curtains, patient cradle, area of iron, gate handles, respiratory concern equipments, tools and procedures implicated throughout wound care, brooms, keyboards etc. [51, 52, 53]. Its discovery and management of spreading is even more complex owed to its imitations, which coexists with extensive strains [54, 55]. *Acinetobacter* colonization is normally originated and noticed in patients who are protected in intensive care surroundings for extended times along with manifold intravenous lines, observing machines, surgical drains, indwelling urinary catheters .etc. in them. Yet no cultural fondness, sexual penchants and forecast for age of this bacteria is present [56].

Motility and morbidity

Mortality features and morbidity shown by *A.baumannii* infection causing strains narrate us the patient's fundamental cardiopulmonary immune position as compared to the intrinsic toxicity of the bacteria. These features and numbers in host showing various infects and illness are amplified mainly because of their own fundamental sickness inspite of the infection cause by this pathogen [56].

Antibiotics resistance mechanisms and methods

Quinolones

Alterations to DNA gyrase or topoisomerase IV throughout transmutations happening the *gyrA* gene as well as *parC* gene has been finely explained for *A. baumannii* [57]. These mutations obstruct the objective site binding. Several other important substrates are also known for multidrug efflux pumps [58], counting the pump AdeABC which is a type of RND [59] along with the AdeM which is a type of MATE pump [60].

Polymyxins

In spite of current reports on growing artificial laboratory built resistance along with hetero-resistance to the polymyxins class [61], machinery of this defense mechanism of *A.baumannii* is indefinite.

β-Lactams

Enzymatic mechanisms:

Nearly all widespread machinery of resistance of *A. baumannii* against β-lactams class is due to enzymatic dilapidation by enzymes known as β-lactamases. Though, in maintaining with the intricate scenery of this organism, several mechanisms frequently work in recital to create the similar phenotype [62].

AmpC cephalosporinases are intrinsically coded by chromosomes of *A. baumannii* strains [63] which are also recognized as *Acinetobacter*-extracted cephalosporinases (ADCs) [64].

Non-enzymatic methods:

Resistance against β-Lactam classes, together with resistance against carbapenem, also contributes to this methodology, containing variations in outer membrane proteins (OMPs) [65], and multidrug efflux channels [66] and variations in the similarity and appearance of penicillin-binding proteins [67].

Aminoglycosides

As described on top, the incidence of genes conventioning for aminoglycoside changing enzymes along with that of type-1 integron are extremely widespread in this pathogen [68]. This rising resistance apparatus damages combining of aminoglycoside to its intentiongenes and proteins and bestows elevated defense towards all such classes of antibiotics [69]. This class of antibiotics also plays the role of important substrates for AbeM pump, which is actually part of the multidrug and toxic amalgam extrusion (MATE) relatives [70].

Tetracyclines and Glycylcyclines

Defence mechanism against this class of antibiotics along with its imitative, are arbitrated through efflux ribosomal resistance [71]. Efflux pumps which are specific to this antibiotic class comprise regions encoded by the *tet(A)* to *tet(E)* predictors, frequently established within gram (-ve) bacteria. Separately from this type of efflux pumps, tetracyclines along with glycylcyclines and their derivatives are vulnerable to efflux shown by various efflux schemes [72].

Remaining classes of Antibiotics

Presence of trimethoprim along with sulfamethoxazole resistance in *A. baumannii* is elevated in lots of regions worldwide [73]. 3' end which is a conserved part of any integron normally have a *qac* gene merged to a *sul* gene, bestowing defence against many antibiotic classes [74]. As a result, resistance shown against sulfonamide has been revealed to be extremely prognostic of most of the isolates of this pathogen [75]. Likewise, resistance against trimethoprim (*dhfr*) and chloramphenicol (*cat*) are coded by some specific genes accounted in genetic structure of *Acinetobacter* [76]. Efflux might also supply to resistance alongside these agents [77].

Biofilm by *A.baumannii*

Biofilm is colonization of microorganisms, mostly bacteria, attaching mutually in a tertiary arrangement and presentation burly observance to the facade. These fanatic cells are habitually well-established within a self-produced environment of extracellular polymeric substance (EPS) in accumulation submitted to as slime which is a polymeric accretion typically gathered of extracellular DNA, proteins, and polysaccharides. Biofilms may form on biotic (epithelial cells, fungal filaments) or abiotic surfaces (polystyrene, glass) and can be unbridled in natural, industrial and hospital environment [78, 79]. *A.baumannii*'s aptitude to continue for extended periods in dissimilar environmental set up acts a major function in its Biofilm arrangement [80, 81]. The purpose of extracellular polymeric matrix is typically to defend the bacteria from environmental conditions, defend from the host and also used as a resistance machinery which usually stimulated throughout the time of infections [82]. It is pragmatic that the solitary clinical isolates of these bacteria which show vulnerability towards disinfectants, antimicrobials, various

environmental stresses etc. were the ones which unite to form Biofilms which in turn confirm to be multi drug resistant and antimicrobial resistant [83]. Biofilm beginning configuration and growth is an extremely synchronized and managed procedure by each of the films planktonic counterparts.

Mechanism of formation

The catalog of the aspects that can allegedly sway the configuration of the Biofilm includes nutrient accessibility, bacterial attachments (pili and flagella), bacterial surface gears (outer membrane proteins, adhesions), and quorum sensing and macromolecular discharges (polysaccharides, nucleic acids) [84]. *A. baumannii* shows adhesion equally to biotic surfaces, w improved through *bla_{PER-1}* gene [85]. Fascinatingly, *A. baumannii* ATCC 17978 strains showed modest biofilm formation on glass exteriors when kept to grow under blue light, whilst usual biofilm creation was pragmatic when kept again in darkness for growth [86]. This observation is arbitrated by the Bls-A photo-receptor protein, which encloses a *N*-terminal blue-light-sensing. It has been stated about varied transcription of *bls-A* at different temperature profiles differentially influences biofilm formation in response to light. Actually it appears to show a worldwide consequence on this pathogens composition, upsetting biofilm structure as well as various other features [86]. Lately, ethanol has also been suggested to affect biofilm configuration on non-living surfaces. In detail, manufacturing of proteins occupied formation of carbohydrates along with lipids was also exposed to augment in ethanol presence, thus increasing biofilm type substance and therefore increasing its biofilm creation and lessening the motility [87]. *Acinetobacter spp.* accounted to create QS-signaling molecules because of which the glueyness and biofilm forming patterns of the bacteria was influenced due to the cell populace [88]. The *A. baumannii* species established *N*-acylhomoserine lactone (*i.e.* *N*-3-hydroxydodecanoyl-homoserine lactone) which is a QS molecule that is significant for biofilm formation on various surfaces [89]. ATCC 19606 strain demonstrated that manufacturing of the pilus like proteins arbitrated by the *CsuA/BABCDE* usher-chaperone congregation scheme was necessary for overall biofilm development [90]. In addition to this, the appearance of the *csuA/ BABCDE* operon was originated to be synchronized by a two-component arrangement composed by the

sensory kinase, which is mainly encoded by *bfmS* and the other one is response regulator which is further encoded by *bfmR*. Suppression of the *bfmS* sensory kinase gene showed a reduction, though not full elimination [91]. A dissimilar contribution of a worldwide transcriptional repressor, which is also a homologue and family member of the histone-like nucleoid structures, have been pragmatic in biofilm creation [92]. Also the outer membrane protein, which is the OmpA, has a key place in the configuration of this pathogen on abiotic surfaces, and in the communication of this pathogen along with other cells [93]. SEM examination of biofilm have revealed that Bap (biofilm-associated protein), at cell surfaces is necessary for three-dimensional tower construction and water channel arrangement on medically applicable surfaces, together with polypropylene, polystyrene, and titanium [94]. Mutagenesis of locus *pglC* banned the fusion of glycoprotein with capsule, consequential in irregular biofilm arrangements. The *pglC* mutant, likewise to the wild kind strain, gave birth to thick cumulates on abiotic surfaces, whereas the study of the biofilm construction exposed an uneven and messy phenotype, signifying a unfocused attachment [95]. Biofilm expansion and maturation by *A. baumannii* clinical isolates depends on poly- β -1,6-*N*-acetylglucosamine (PNAG) production which is prearranged by a group of four genes (*pgaABCD*) [96].

Aims and Objectives:

1. To check the biofilm forming capacity of *A. baumannii* isolates from different clinical sources (sputum, pus and peritoneal fluid).
2. To compare the antibiotic resistance of the isolates with their biofilm forming capacity along with their clinical sources (sputum, pus and peritoneal fluid).
3. To compare the twitching motility phenotype of the isolates with their clinical sources (sputum, pus and peritoneal fluid) and biofilm forming capacity
4. To check for the viability of the usage of copper nanoparticles also copper nanoparticles along with amalgam of catharanthus-rosesus plant extract, as an antibiofilm compound.

Chapter 3: Materials and methods

***A.baumannii* identification and validation by colony PCR method:**

PCR assays present the prospective for fast discovery and species recognition of pathogens [97]. A colony PCR-based assay was used for this project that can identify *A. baumannii*. *Acinetobacter* species recognition was done by a spacer region called as ITS region [98]. Primers ITS-F (59-CATTATCACGGT-AATTAGTG-39) along with ITS-R (59-AGAGCACTGTGCACTTAAG-39) were made in use for amplifying an portion of this mentioned region of size of around 208 bp. For the PCR assays of the *A.baumannii* (ATCC 19606 AND 1605) isolates, the DNA template was all set by boiling technique [99]. Momentarily, single colony of an untainted and pure culture was balanced in 20 μ l of nuclease free water and heated at 95 $^{\circ}$ C for 10 min in a thermocycler machine. Following centrifugation in a micro centrifuge (bench top lab systems) at 6000g for three minutes, after which the supernatant collected was kept at -20 $^{\circ}$ C. PCR was performed with 2 μ l of template DNA (in duplicates) in a total reaction volume of 5 μ l consisting of 7.5 μ l of master mix, 0.5 μ l (F and R both) primer and 4.5 μ l of nuclease free water to make up the total volume. The PCR program consisted of an initial denaturation step done at 94 $^{\circ}$ C for 5 min, which was repeated for about 35 cycles of final denaturation done at 94 $^{\circ}$ C for 1 minute, annealing was performed at 58 $^{\circ}$ C for 30 sec time, and an initial extension which was kept at 72 $^{\circ}$ C for 30 sec time, with a final extension step as the last step at 72 $^{\circ}$ C for 5 minutes. A thermocycler PCR was used.

Table 1: Reconstitution (1:10 dilution) of lyophilized primers:

Primer	Yield(nano-moles)	Working volume(μ l)
ITS-F	40.7	407
ITS-R	25.7	257

The primers was synthesized and lyophilized by EUROFINS genomics for this project.

PCR gel electrophoresis:

The PCR amplified products with the ITS gene primers were seen and verified on 1.5% Agarose gel in 150 ml of 1X TAE buffer. To which 10µl of ETBR was added. 15µl of the above gained PCR samples was used in the well, run with a 100bp ladder (NEB) as a reference and standard.

Biofilm formation and quantification:

In totality 48 *A. baumannii* strains were gathered from Department of Microbiology (Bacteriology division), AIIMS, New Delhi during the examining stage. Clinical sources of isolation of *A. baumannii* incorporated patient's pus, respiratory and peritoneal fluid. Assembled strains were typified by typical biochemical assays (carbohydrate and amino acid utilization). Two standard strains of *A. baumannii* (ATCC 19606 and ATCC 1605) was also used as controls for antimicrobial susceptibility examination and biofilm creation assay [100].

Biofilm formation in *A. baumannii* is to be studied on two strictures:

- Time dependent quantification of biofilm (24 hours & 48 hours) {4 plates}.
- Temperature dependent quantification of biofilm (24 hours at 37°C and 44°C, 48 hours at 37°C and 44°C) {8 plates}.

Culture Preparation:

Every *A.baumannii* strain was matured independently in Luria-Bertani medium (LB) overnight at 37°C and every sample from the above medium was again freshly-cultured for about 3X in LB for a time period of 24 h at 37° C along with shaking (200 rpm). These required cells were produced from above mentioned overnight grown bacterial cultures by the means of centrifugation performed at 3,600 g for 30 min at 4° C. Followed by washing of cells, which is performed 2 times in sterile 1x PBS [pH 7.2] and the bacterial cell pellet was then re-suspended in PBS again, which was further used as the inoculums. For the corroboration of the bacterial populace in the inoculums, McFarland Nephelometer Standards No. 0.5 was made and used [101]. Two hundred micro liters of the washed culture at 0.2 O.D. were used as the inoculums (6.0 logs CFU) (O.D. 0.1). Inoculation of sterile 96-wells polystyrene plates with 200 microL of each above prepared bacterial cell suspension (6.0 log CFU) was done.

Following incubation of the mentioned prepared culture plates at 37° C and 44° C for 24 and 48 hours, which is devoid of any agitation for biofilm production, the planktonic cells stuck on the sides of plate is washed properly, and with rigor, with already prepared PBS solution to detach these cells from the walls which was followed by keeping the plate upside down to dry followed by staining step with freshly prepared and filtered 1% crystal violet stain and kept for 15 min. The wells were washed once again with 200µl of freshly prepared ethanol-acetone solution (80:20 v/v) to solubilize the crystal violet dye complex and the biofilm produced in the wells. The optical density is then preferred to be taken at 570 nm (OD 570) which was determined using BIORAD spectrophotometer. Each step was performed three times and the average optical density was calculated for further use [102].

The following values were allocated for biofilm determination:

* Classify isolates as biofilm-forming if they capitulated the OD values that were at least twice those of the negative controls.

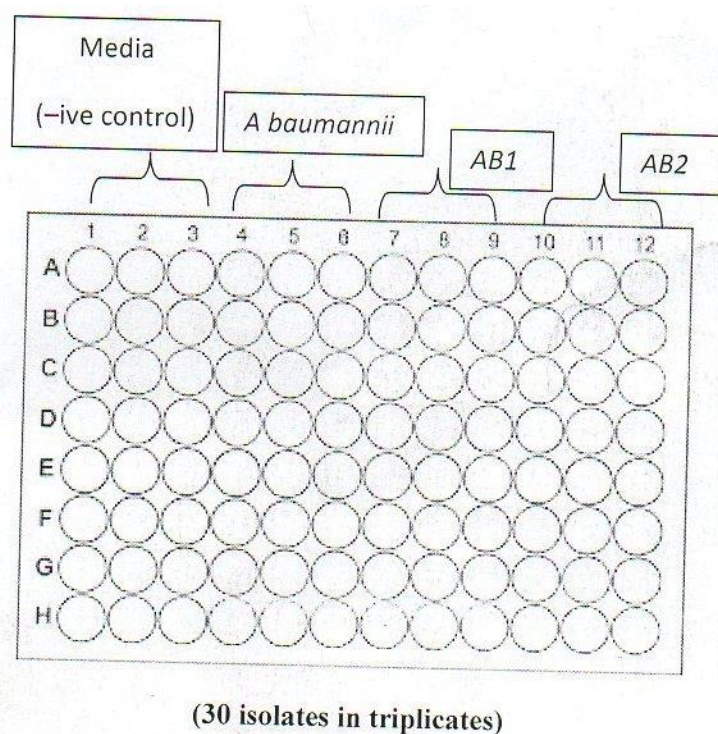
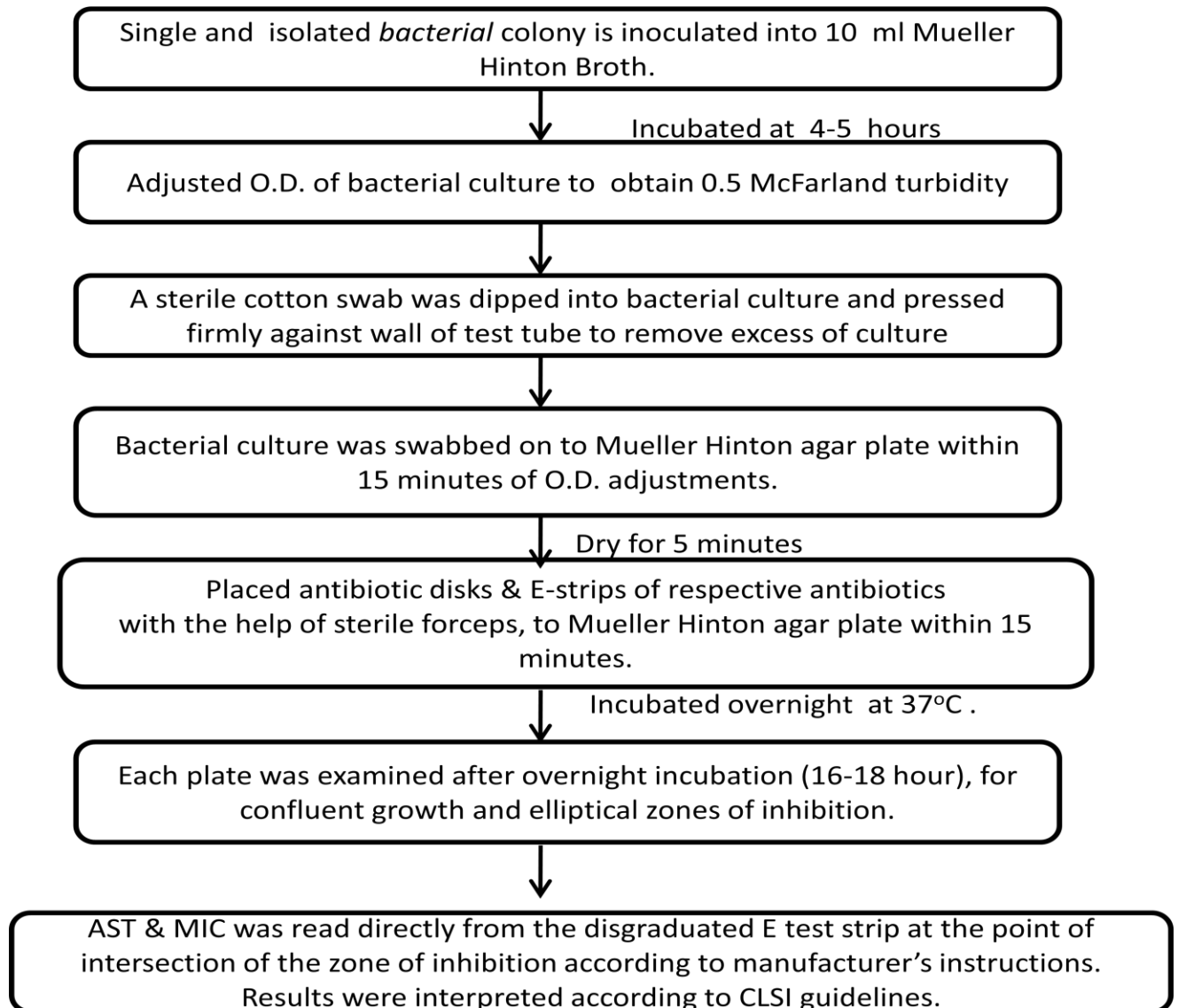


Fig 1: 96-well plate with allocation of isolates in triplicates

Biofilm creation capability in *A. baumannii* strains was also observed under bright field microscopy. Microscopic analysis also holds and authenticates the different capacities of biofilm producing isolates, qualitatively.

Antibiotic susceptibility test (AST) and minimum inhibitory concentrations determination (MIC):



Antibiotics were chosen according to ICMR guidelines: piperacillin tazobactam, cefepime, levofloxacin, tetracycline, amikacin, imipenem, meropenem, cefoperazone sulbactam, ceftazidime [103].

Motility Assay:

Modified LB broth in the company of 0.4% w/v agar was made in use, for the motility assays done in the project for *A.baumannii* strains. A single isolated colony of bacterium is inoculated into 10 ml Luria broth at 37°C for overnight. Newly grown cultures were pierced to allow multiply and spreading of bacteria on the surface of the medium (which is 0.4% semisolid as mentioned above) that is only done for swarming type of motility and at the junction between the base of the Petriplate and L.B-medium (which is 0.8% semisolid) which is performed for twitching motility [104]. Plates were equipped on the similar day as on which inoculations are done. Subsequent to the above procedure, the plates were covered properly with parafilm and further incubated at 37°C for 48 hours.

For elucidation of isolates showing positive results, were announced to be the strains which demonstrated a region of >10 mm about the location of inoculation. For the twitching type of motility, after the inoculation, the agar was useless and thrown away, and the plates were tainted with freshly prepared 0.2% crystal violet previous to apparition and snapped. Assays were executed at least 3X, for every possible isolate for confirming the results [105].

Antibiofilm activity of nanoparticles:

Antibiofilm activity of two nanoparticle solutions was performed by using a standard *A. baumannii* isolate, namely *A. baumannii* ATCC 1605 (multidrug resistant). Two nanoparticle solutions were taken in the study: sample 1 was nanoparticles synthesized from copper, plant extract and potassium iodide (KI) (0.5 M), sample 2 was nanoparticle made with copper combined only with potassium iodide (0.5 M). From this concentration working dilutions were made of 2mM, 1mM and 0.5mM for the test sample. Plain culture of ATCC 1605 was taken as negative control in a 96-well plate. Then the nanoparticles along with calculated volume of cultures to sum to 100µl are placed in the 96-well culture plates. After which plates were kept for incubation for biofilm production and their destruction due to antibiofilm particles, at 37°C for the required time profiles. At the completion of incubation the plates were given 1X PBS

washing, twice, followed by staining each well with 1% crystal violet and keeping the stain for 30 minutes after which again single washing of PBS was given.

Biofilm formation restriction property of the above mentioned samples was checked by performing light microscopy and imaging of each well and plate. This was followed by dissolving the cells and biofilm with ethanol-acetone (80:20 v/v) solution and the O.D of the plates were taken at 570nm and recorded, to quantify the biofilm formation.

Table 2: required concentrations of nanoparticle sample 1 as well as for sample 2 with required *A.baumannii* (for ATCC 1605) culture volumes.

Molar concentration(mM)	Nanoparticles(μ l)	Culture(μ l)
2	0 (negative control)	100
2	51.2	48.8
2	25.6	74.4
2	12.8	87.2
2	6.4	93.6
2	3.2	96.8
2	1.6	98.4
2	0.8	99.2
2	0.4	99.6
1	0.2	99.8
0.5	0.2	99.8

The antibiofilm activity was studied for four different time profiles .i.e. is for 2 hours, 4 hours, 6 hours and 8 hours.

Chapter 4: Results

Colony boiling PCR method:

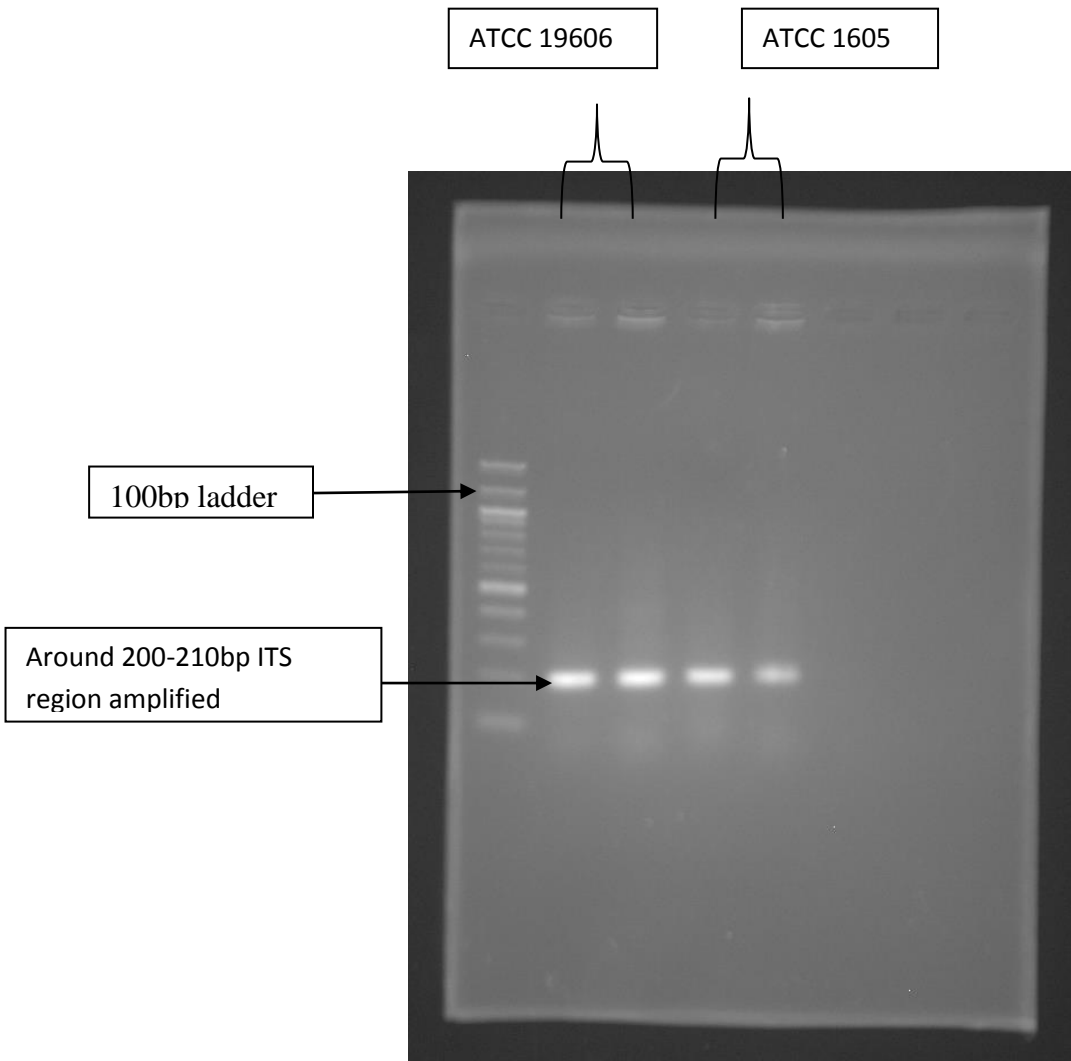
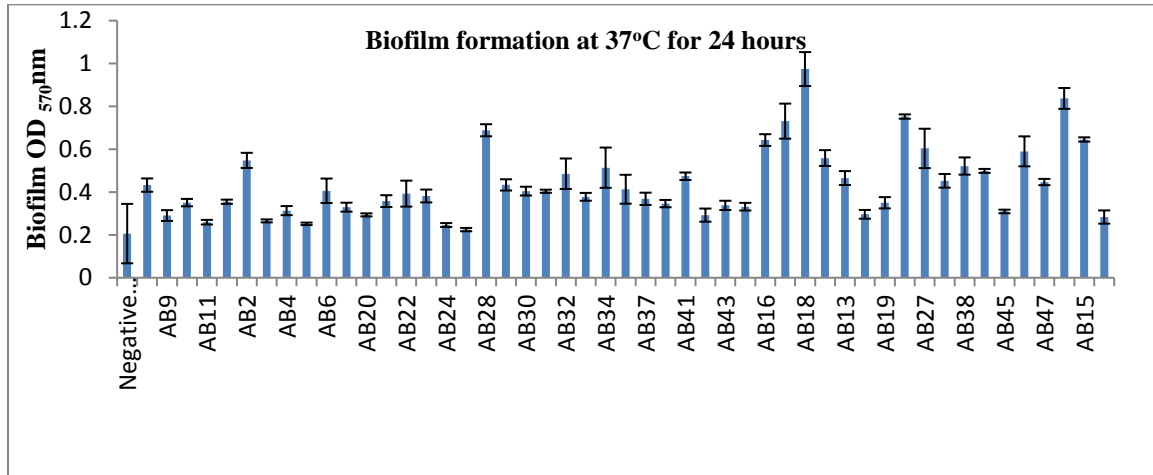


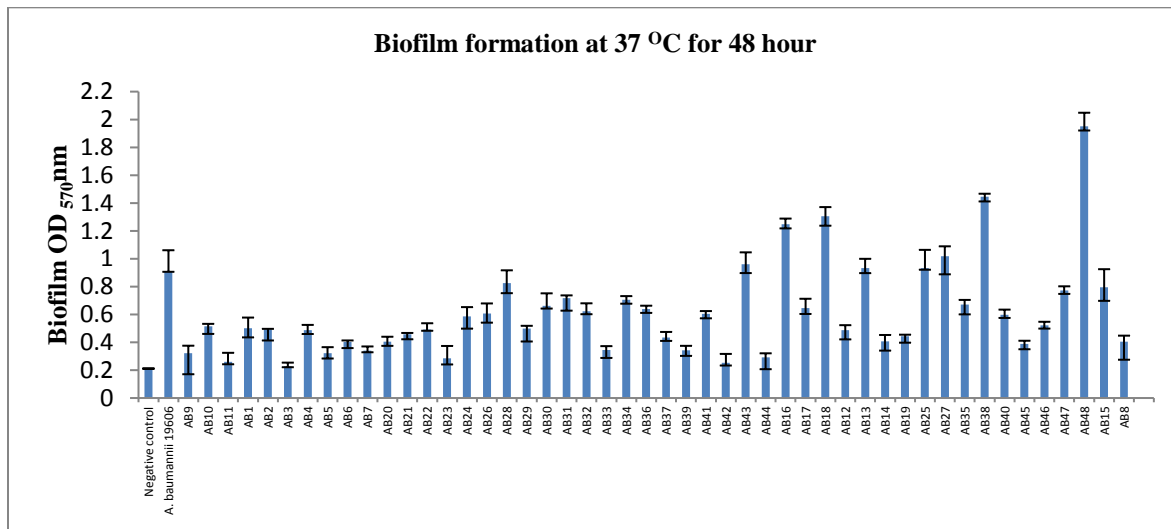
Fig 2: PCR gel electrophoresis with amplified ITS region bands of around 200bp size.

Biofilm formation and quantification:

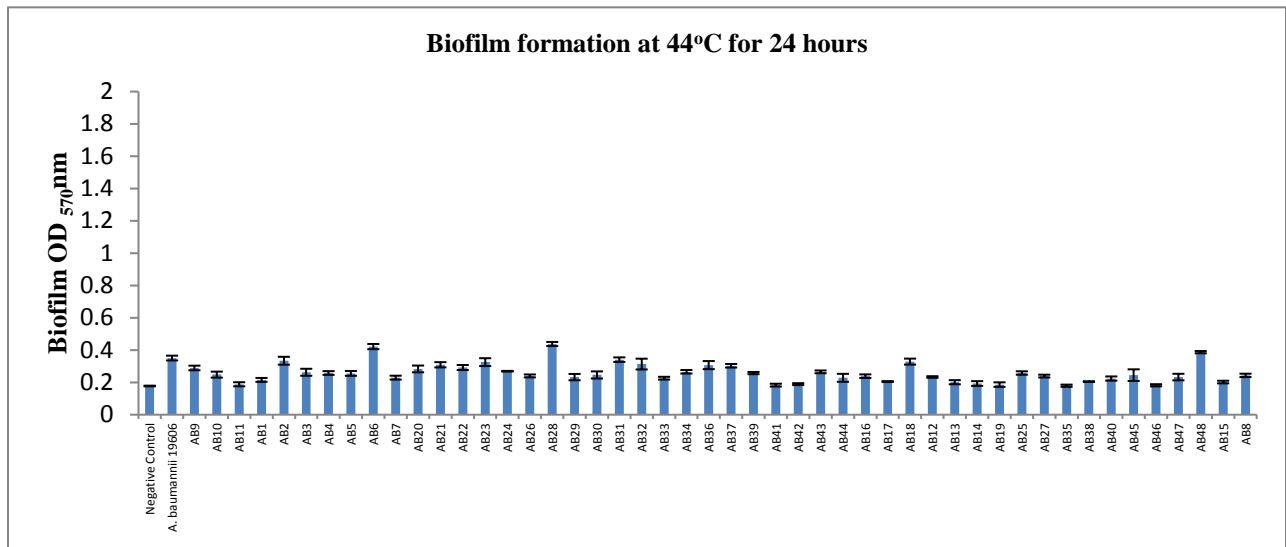
The graphs after plotting the observed O.D for the mean, standard deviation and standard error for triplicates of each strain and sample were:



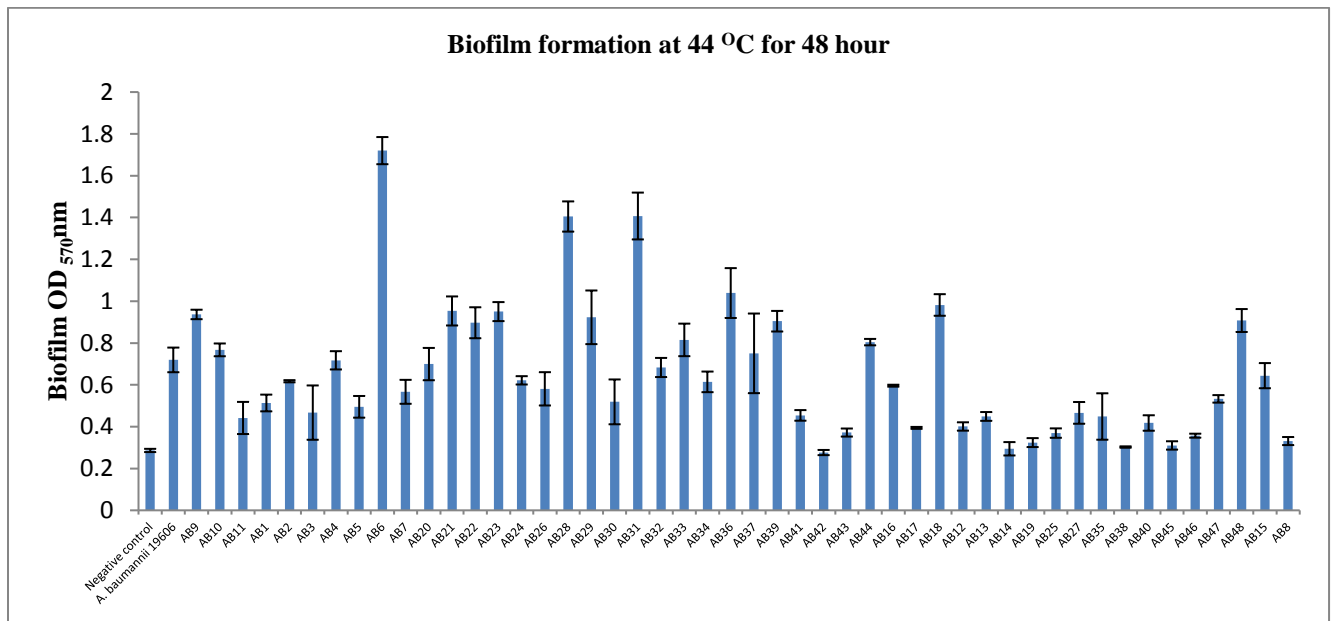
(A)



(B)



(C)

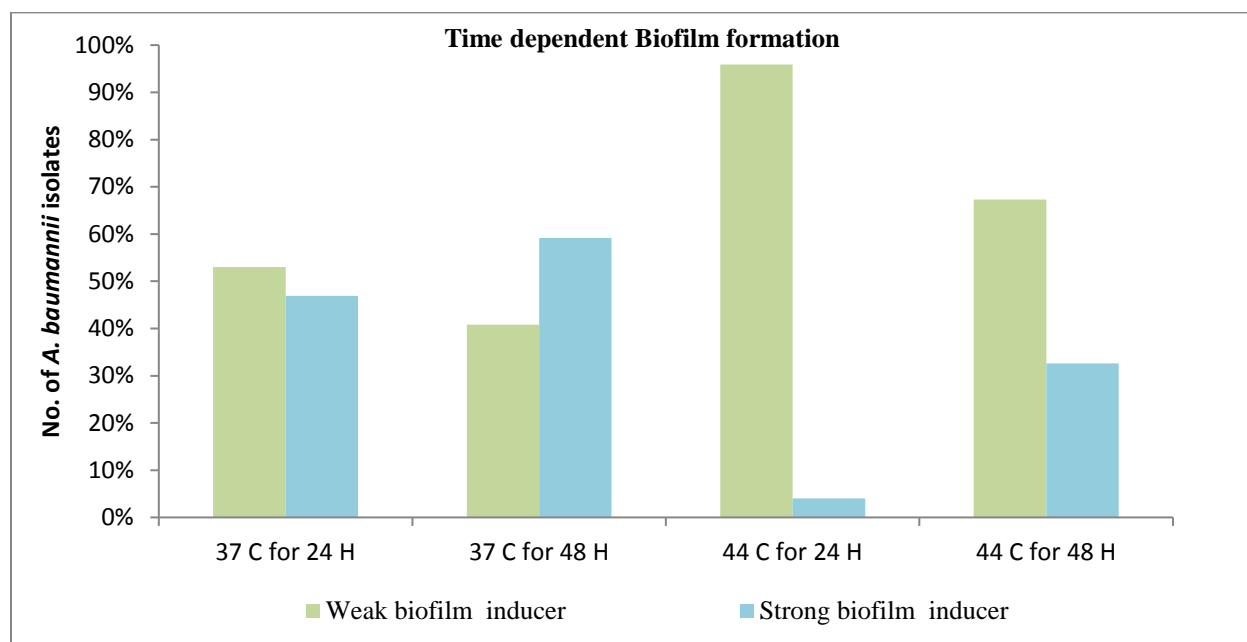


(D)

Fig 3: A, B, C and D: Quantification of biofilm mass by crystal violet assay. Bar diagram shows OD₅₇₀ mean \pm SE (y-axis) against different *A. baumannii* isolates on (x-axis). Each value represents the triplicate of experiments. Panel A& B represents biofilms production at 37°C (24 h and 48 hours) while C&D shows the biofilm expression at 44°C for 24 and 48 hours.

	No of <i>A. baumannii</i> Strains			
	37 C for 24 H	37 C for 48 H	44 C for 24 H	44 C for 48 H
Weak biofilm inducer	53%	40.80%	95.90%	67.30%
Strong biofilm inducer	46.90%	59.18%	4%	32.60%

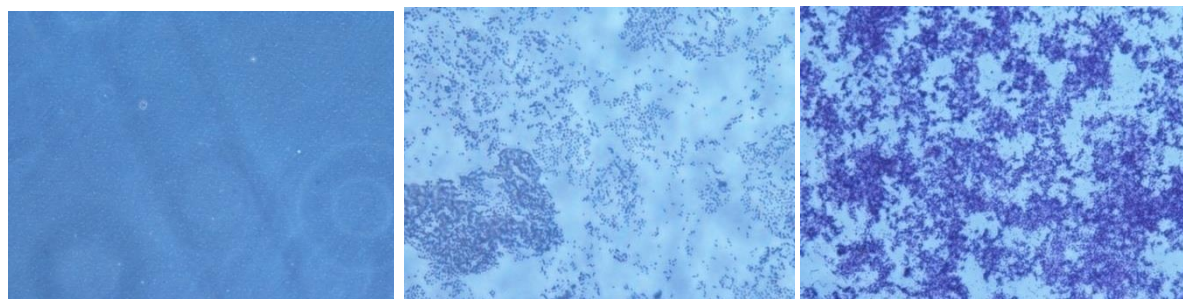
Table 3: Comparison of percentages of weak and strong biofilm formers with respect to time.



(E)

Fig 4: image E is Time dependent analysis of biofilm producing ability of isolates of *Acinetobacter baumannii*.

Ability to form biofilm at elevated temperature combined with multidrug resistance, might contribute to the survival of these organisms and their dissemination in the hospital environment.

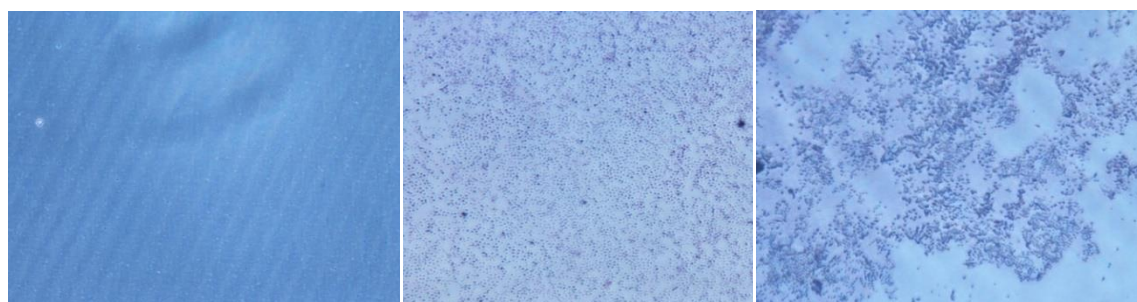


Negative

37°C for 24 h

37°C for 48 h

Panel A.



Negative

44°C for 24 h

44°C for 48 h

Panel B.

Fig 5: Panel A and B are time and temperature dependent bright field microscopy analysis (400X) of *A. baumannii* isolates attached to polystyrene surfaces with crystal violet staining (1% w/v for 10 minutes). Panel (A) represents biofilms grown at 37°C for 24 and 48h and negative control while, panel (B) shows biofilms grown at 44°C for 24 and 48h and negative control.

AST and MIC results:

Table 4: Antimicrobial susceptibility patterns of *A. baumannii* isolated from different sources.

Source	Antibiotics	PTZ	LEV	TE	CPM	CAZ	CPS	MRP	IPM	AMK
Respiratory	S	4%	8%	20%	8%	16%	8%	12%	4%	24%
	R	81%	78%	78%	81%	81%	74%	67%	85%	78%
	I	16%	12%	4%	12%	4%	20%	20%	8%	0%
Pus	S	33%	0%	33%	0%	0%	33%	33%	0%	33%
	R	33%	100%	33%	100%	100%	33%	33%	100%	67%
	I	33%	0%	33%	0%	0%	33%	33%	0%	0%
Peritoneal fluid	S	50%	50%	50%	50%	50%	50%	50%	50%	50%
	R	50%	50%	50%	50%	50%	50%	50%	50%	50%
	I	0%	0%	0%	0%	0%	0%	0%	0%	0%

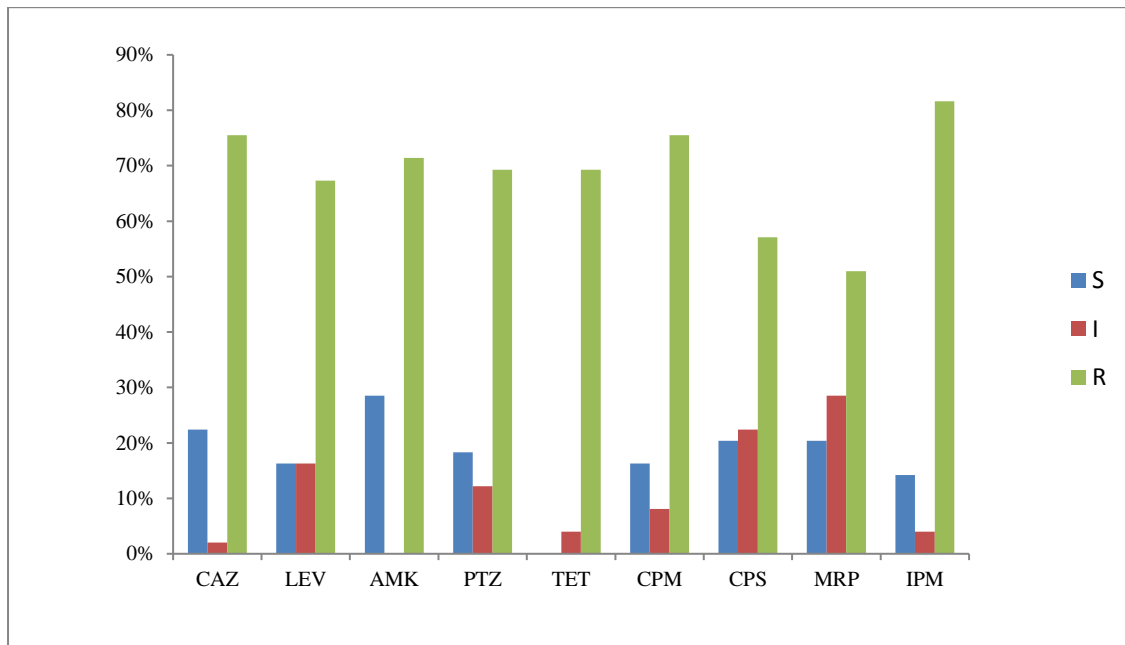
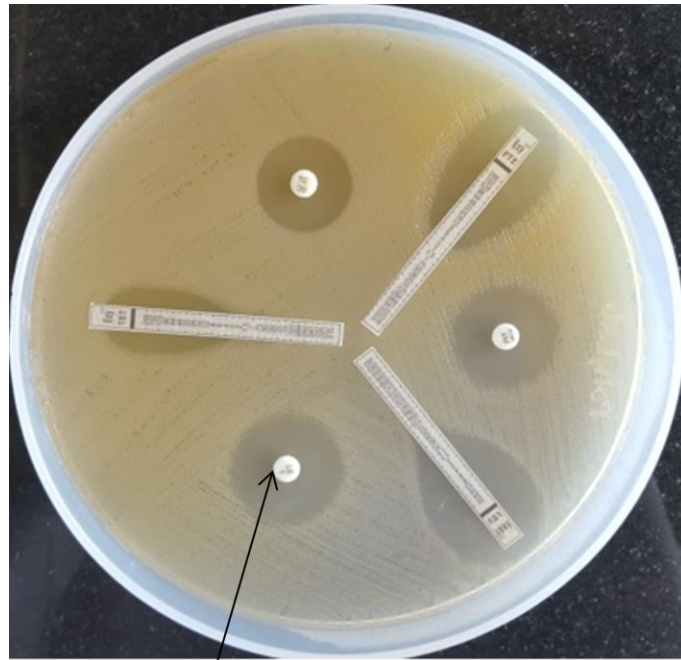


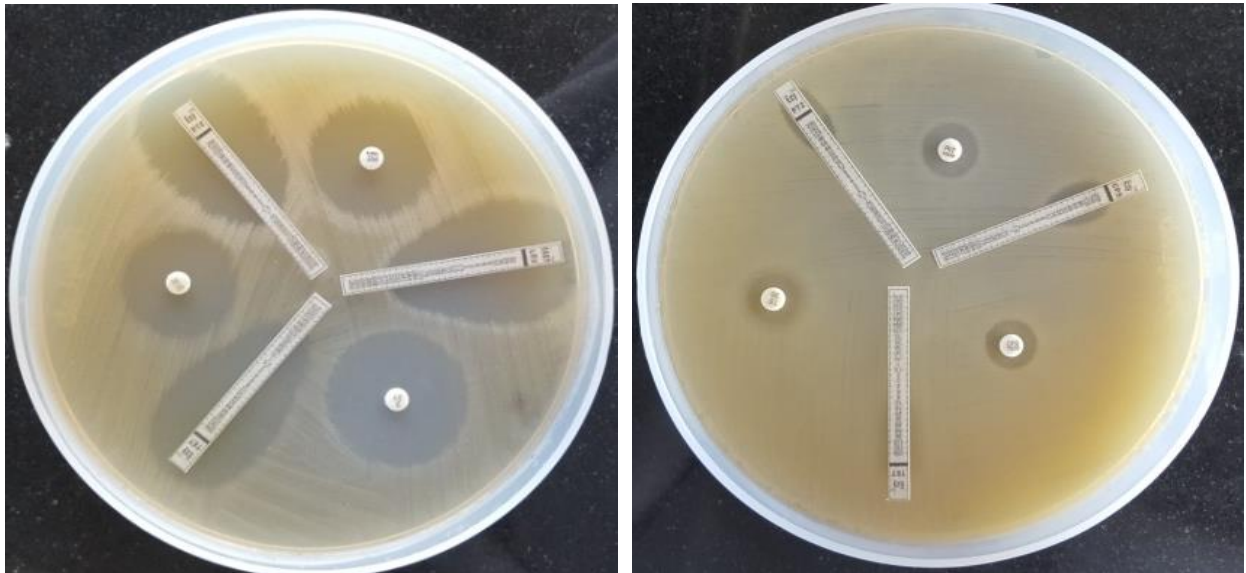
Fig 6: Antibiotic resistance phenotypes of *A. baumannii* isolates. X-axis represents different *A. baumannii* isolates and Y- axis represents percentage of isolates S=Susceptible, I=Intermediate and R= Resistant.

CAZ –ceftazidime, LEV-levofloxacin, AMK- amikacin, PTZ- piperacillin tazobactam, TET-tetracycline, CPM-cefepime, CPS-cefoperazone sulbactam, MRP- meropenem, IPM- imipenem.

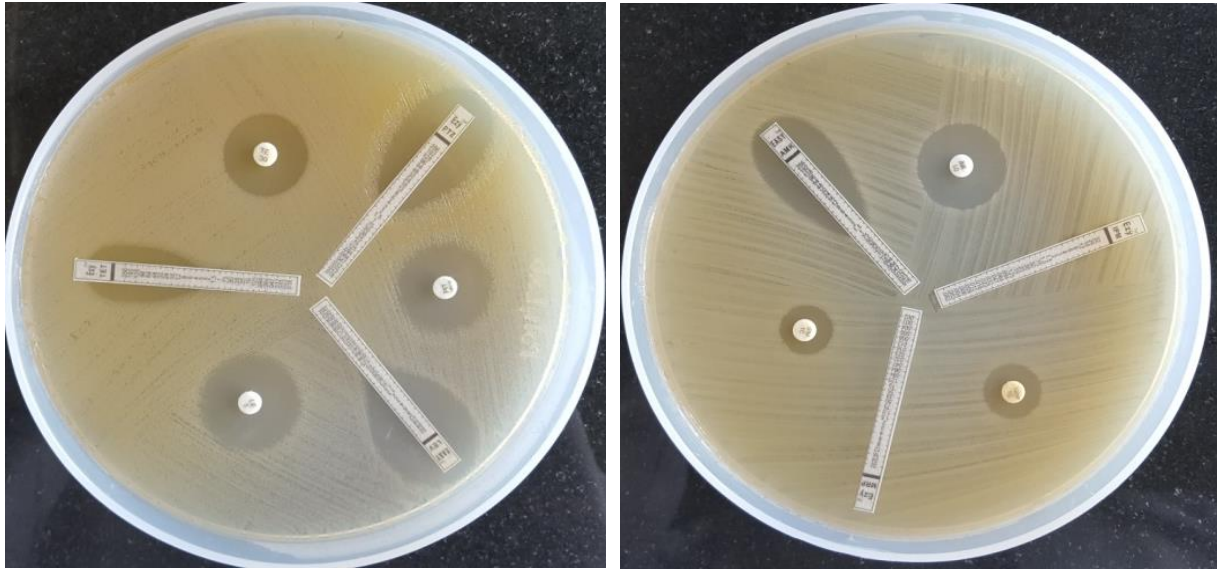


(A)

Zone of lysis



(B)



(C)

Fig 7: Comparative analysis of antimicrobial susceptibility patterns and minimum inhibitory concentrations of *Acinetobacter baumannii*

(A) Type strain ATCC 19606;

(B) Drug susceptible strain AB41 & drug resistant strains AB32 from Respiratory isolates.

(C) Drug susceptible AB17 and drug resistant AB18 strains from Pus isolates.

Motility assay:

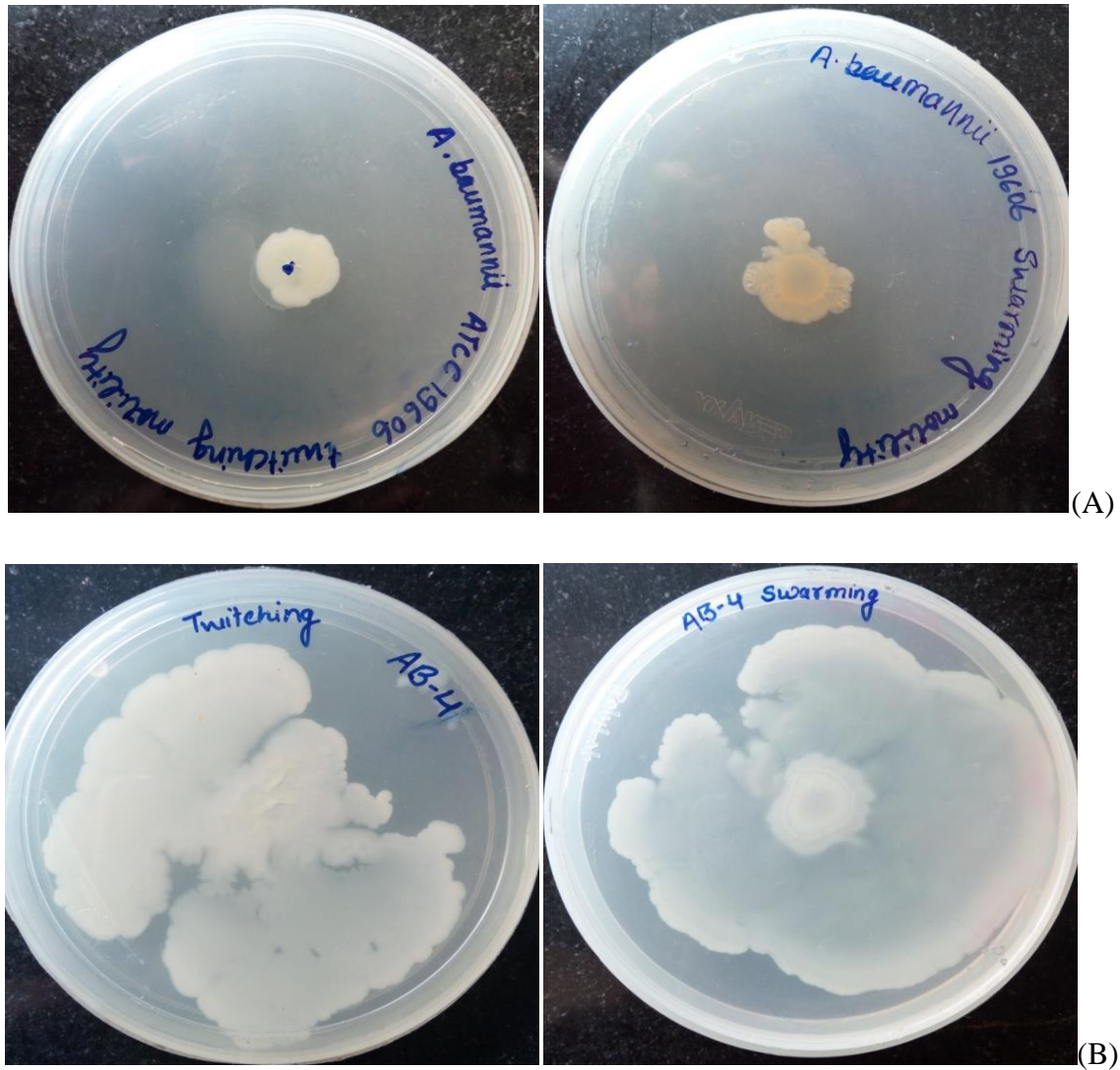


Fig 8: Representing twitching and swarming motility assays performed in semisolid 0.4% modified LB agar.

Panel A: represent non motile multidrug resistant control strain *A. baumannii* ATCC 1605.

Panel B: represent *motile A. baumannii* isolate from respiratory source.

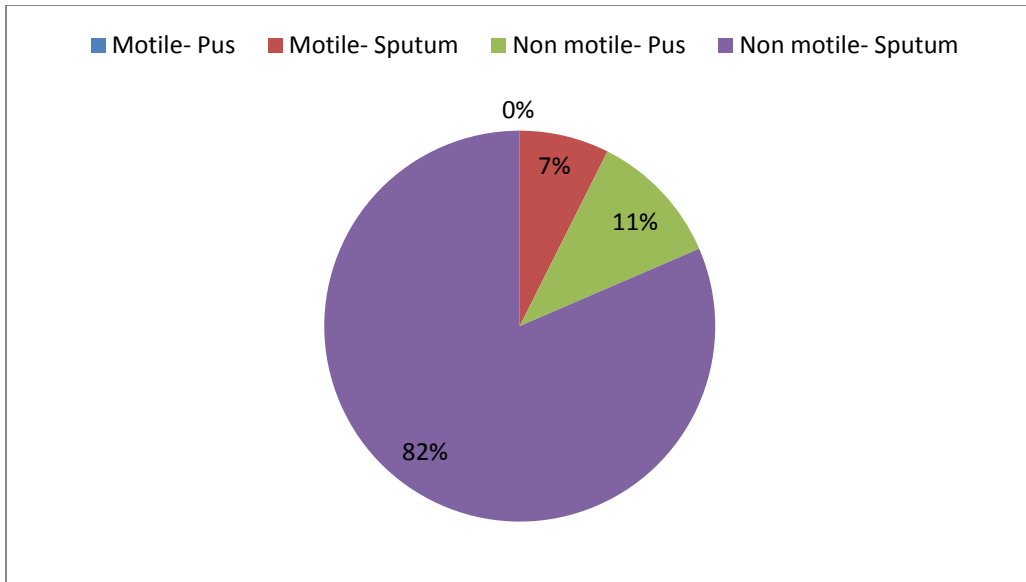


Fig 9: pie chart showing comparison of motility percentages of *A.baumannii* isolates of pus and sputum sources post-staining.

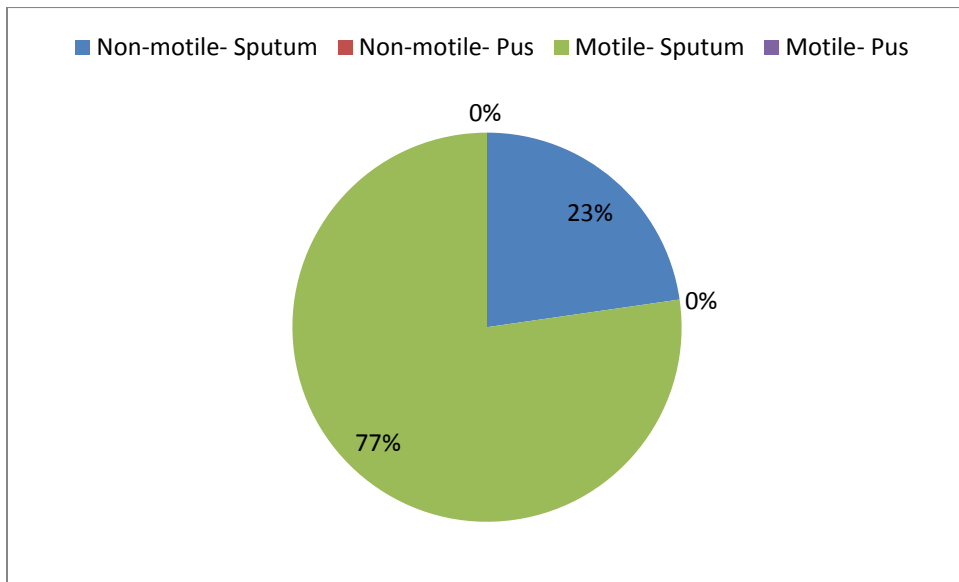


Fig 10: comparison of motility of different *A.baumannii* isolates from different sources before staining step.

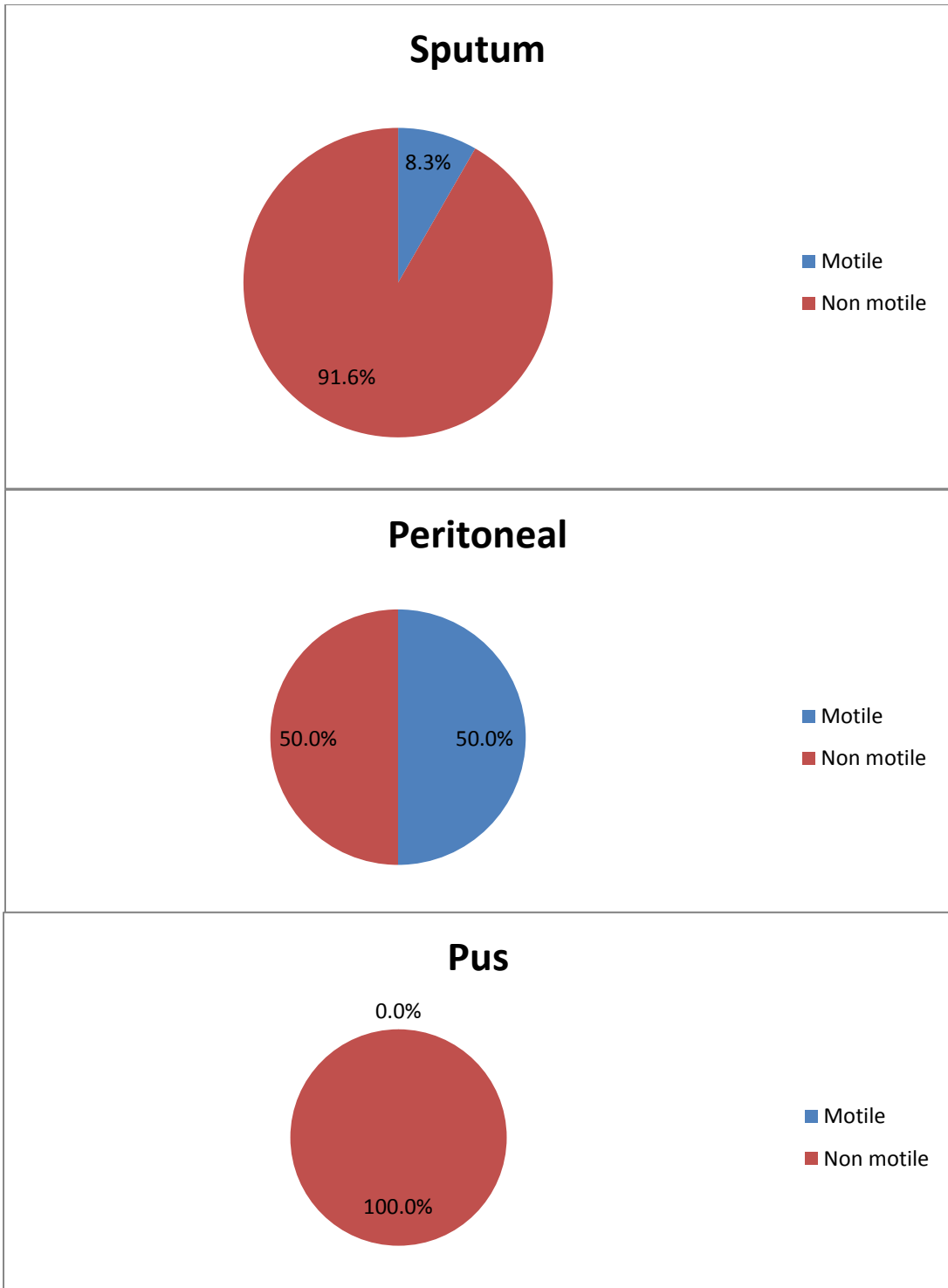


Fig 11: individual pie charts showing motility of isolates from various sources.

sources	Motile (%)	Non-motile (%)
sputum	8.3	91.6
pus	0	100
Peritoneal fluid	50	50

Table 5: percentage representation of motility results of different sources isolates.

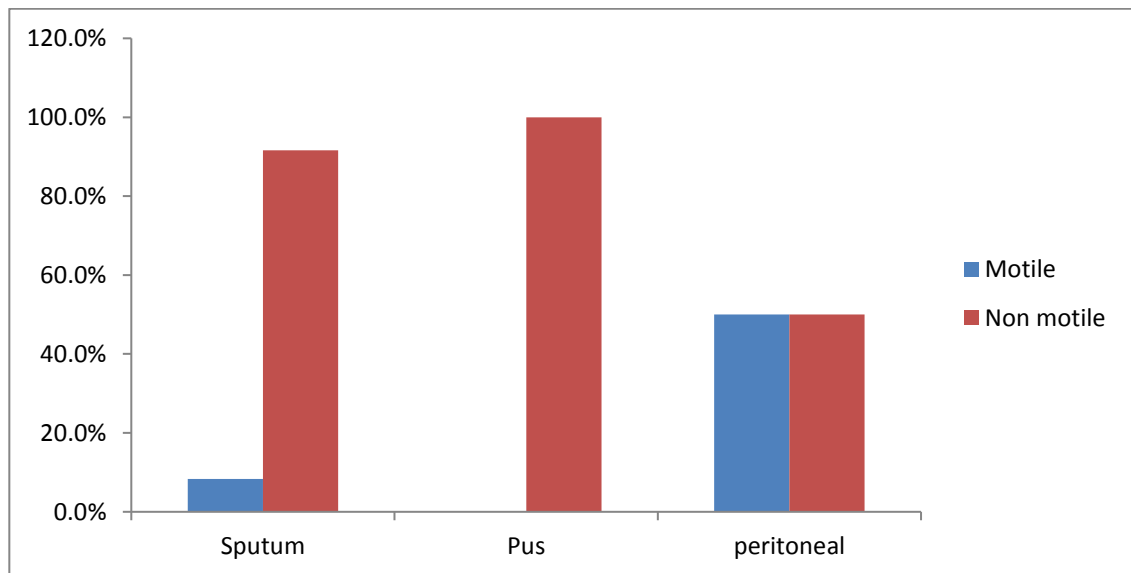


Fig 12: comparison of motility of different sources isolates of *A.baumannii* in single bar graph

Antibiofilm assay:

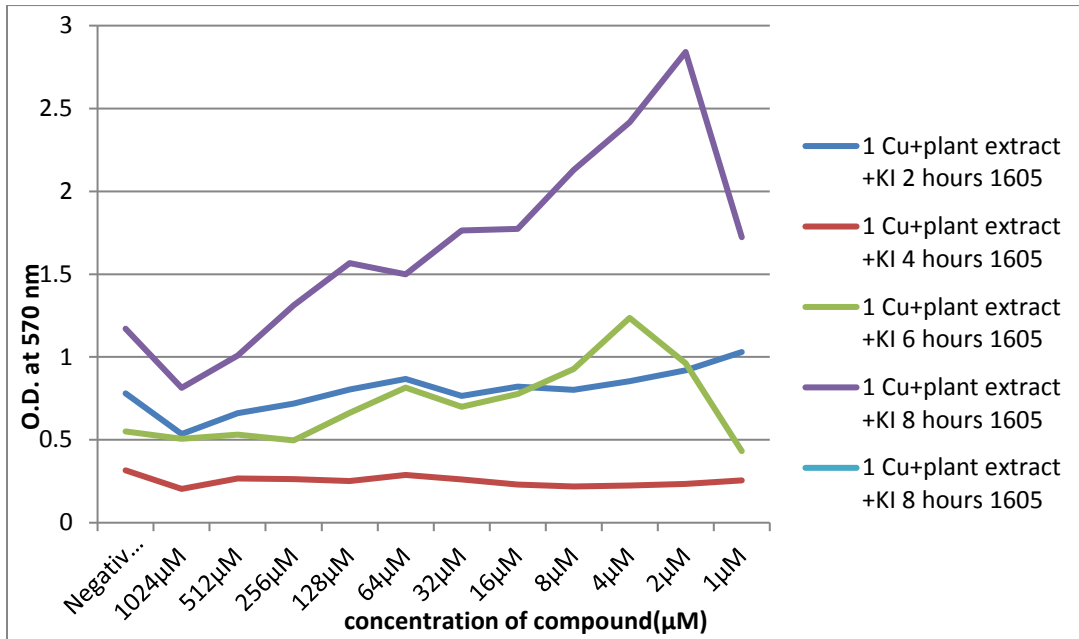


Fig 13: quantification of antibiofilm assay of sample 1(copper + catharanthus-rosesus plant extract + potassium iodide) at three variant temperate profiles i.e. at 2hours, 4hours, 8hours.

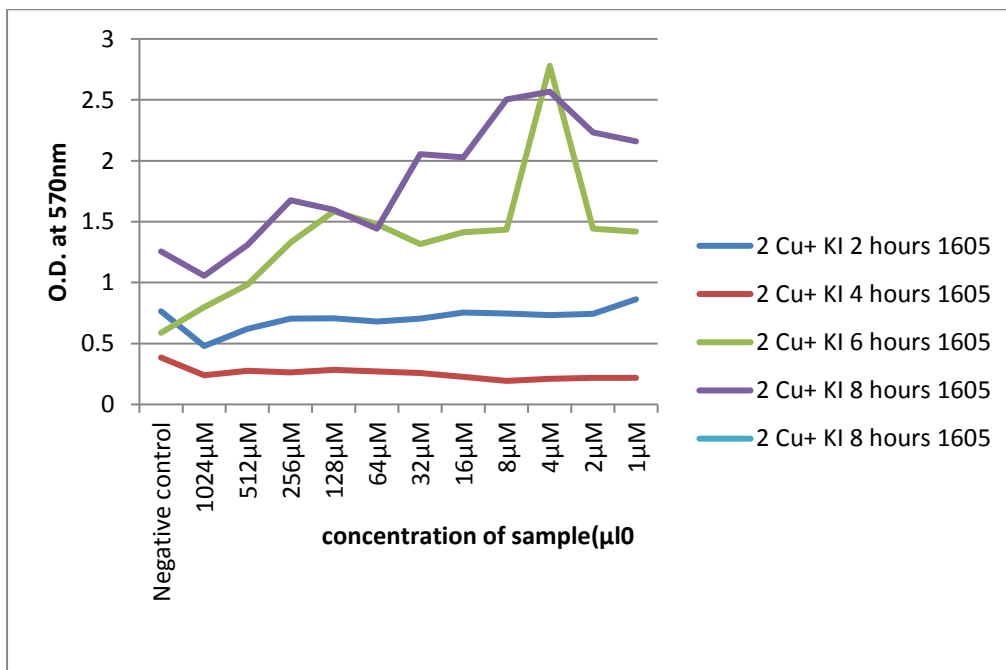
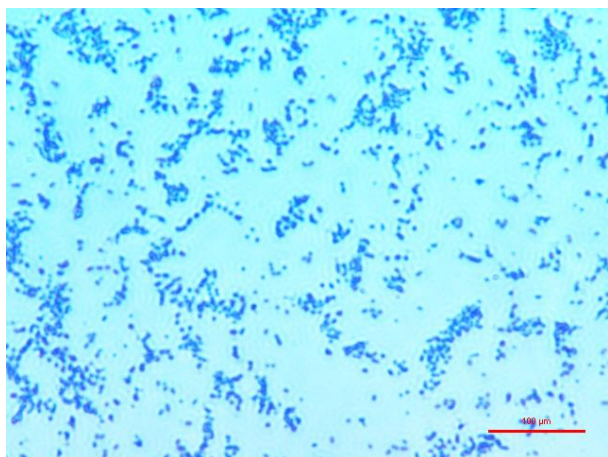
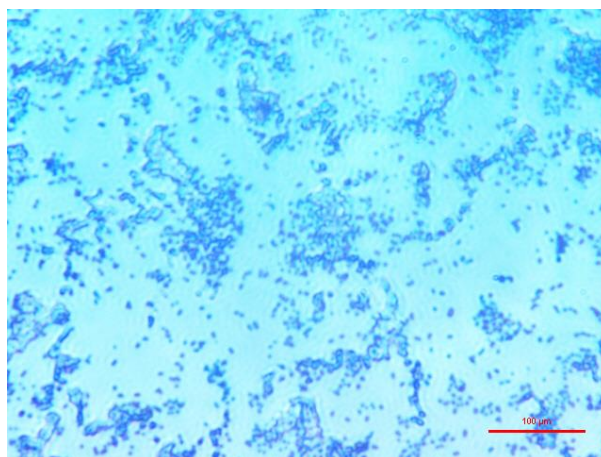


Fig 14: quantification of antibiofilm assay of sample 1(copper + potassium iodide) at three variant temperate profiles i.e. at 2hours, 4hours, 8hours.

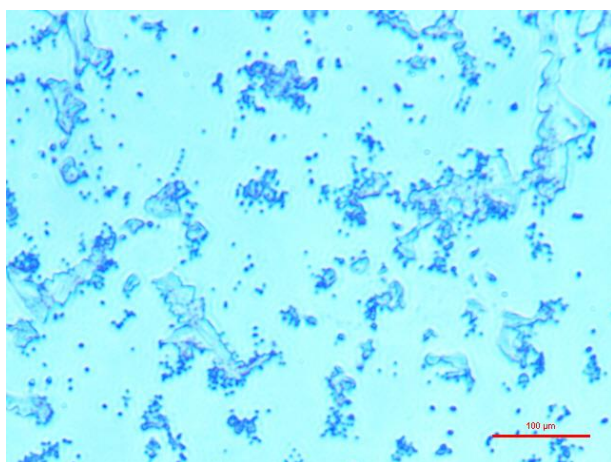


(A)

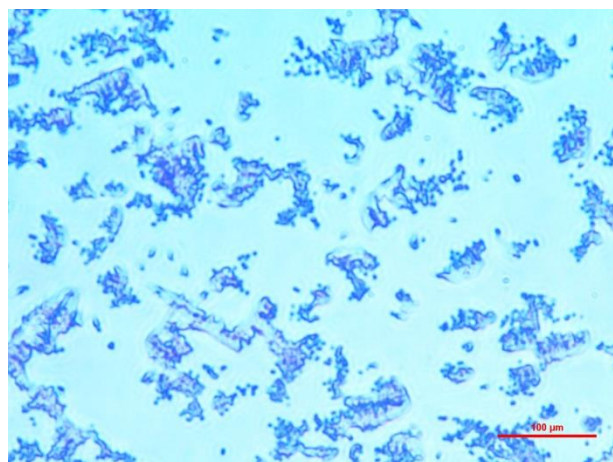


(B)

Fig 15: A and B are time dependent bright field microscopy analysis (400X) of antibiofilm activity of sample-2 against *A. baumannii* isolates attached to polystyrene surfaces with crystal violet staining (1% w/v for 30 minutes) for 2 hours. Panel (A) represents biofilm formation at 1024 μ M concentration of sample 2 while, panel (B) shows biofilm formation at 256 μ M conc. of the sample 2.

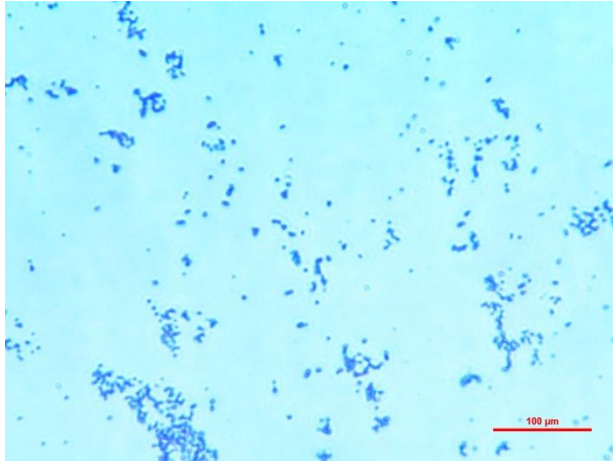


(C)

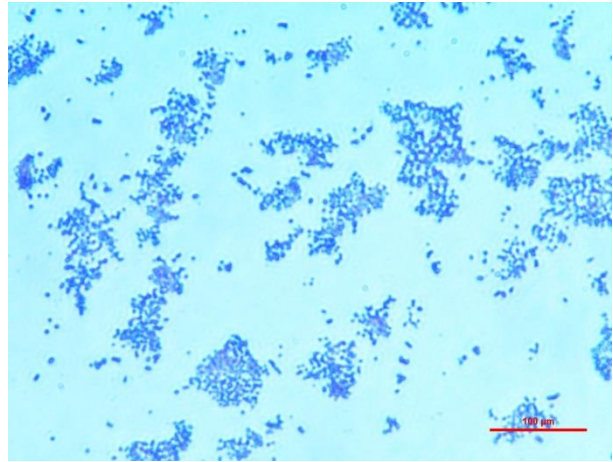


(D)

Fig 16: C and D are time dependent bright field microscopy analysis (400X) of antibiofilm activity of sample-1 against *A. baumannii* isolates attached to polystyrene surfaces with crystal violet staining (1% w/v for 30 minutes) for 4 hours. Panel (C) represents biofilm formation at 1024 μ M concentration of sample 1 while; panel (D) shows biofilm formation at 0.5 μ M conc. of the sample 1.

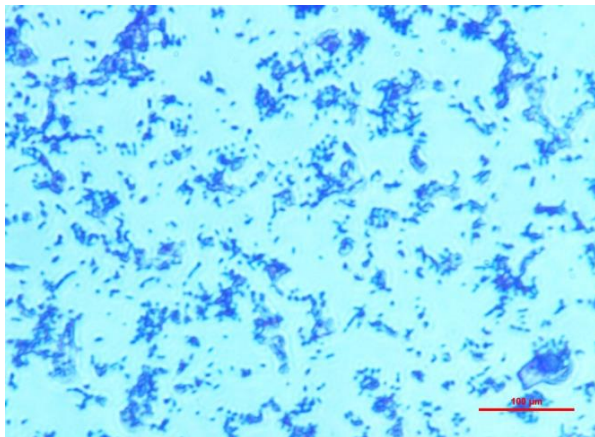


(E)

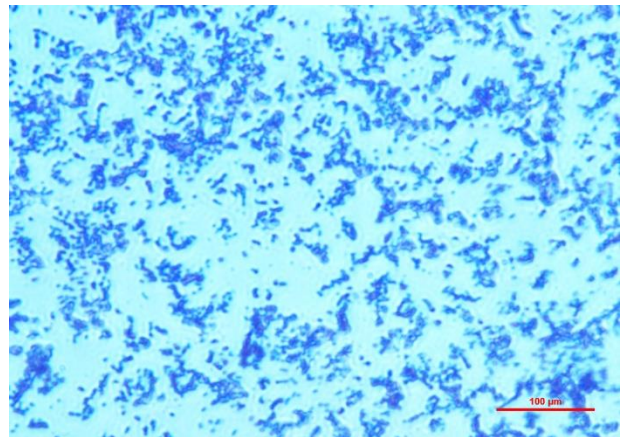


(F)

Fig 17: E and F are time dependent bright field microscopy analysis (400X) of antibiofilm activity of sample-2 against *A. baumannii* isolates attached to polystyrene surfaces with crystal violet staining (1% w/v for 30 minutes) for 4 hours. Panel (E) represents biofilm formation at 256 μ M concentration of sample 2 while, panel (F) shows biofilm formation at 32 μ M conc. of the sample 2.

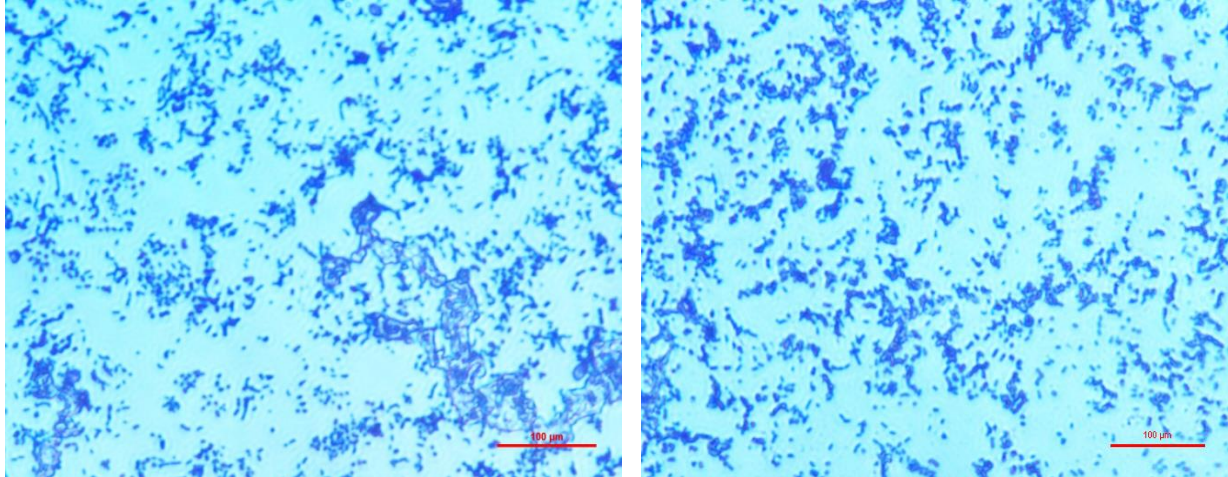


(G)



(H)

Fig 18: G and H are time dependent bright field microscopy analysis (400X) of antibiofilm activity of sample-1 against *A. baumannii* isolates attached to polystyrene surfaces with crystal violet staining (1% w/v for 30 minutes) for 8 hours. Panel (G) represents biofilm formation at 1024 μ M concentration of sample 1 while; panel (H) shows biofilm formation at 0.5 μ M conc. of the sample 1.



(I)

(J)

Fig 19: I and J are time dependent bright field microscopy analysis (400X) of antibiofilm activity of sample-2 against *A. baumannii* isolates attached to polystyrene surfaces with crystal violet staining (1% w/v for 30 minutes) for 8 hours. Panel (I) represents biofilm formation at 1024µM concentration of sample 2 while, panel (J) shows biofilm formation at 0.5µM conc. of the sample 2.

Chapter 5: Discussion

Colony boiling PCR validation:

The ITS region is a conserved region used for identification of *A.baumannii* [d3], as these ITS genes have lower rates of variations in different strains and are mostly the conserved regions having housekeeping genes.

Preciseness of this ITS region amplification which is used for identification of the *A. baumannii* strains was further assured by Agarose gel electrophoresis which showed amplified bands of the *A.baumannii* ATCC 19606 and 1605 after using the specific ITS region primers. The gel picture viewed under GELDOC verified the results of PCR.

Biofilm formation and quantification:

Our results showed that 37°C is more appropriate temperature for biofilm formation as compared to 44°C. Mature biofilm formation occurred at prolonged incubation periods i.e. at 48 hours. However at elevated temperature conditions (44°C) more than 30% *A. baumannii* isolates were capable of forming mature biofilms following incubation for 48 hours.

MDR isolates of *A. baumannii* from respiratory and pus showed higher biofilm formation at 37°C for 24 hours to 48 hours. However, there was no significant association found between susceptible phenotype with high production of biofilm.

At 24 hours of incubation period, >50% of isolates produced biofilm at 37°C, while, few isolates produced biofilm at 44°C (~4%). After prolonged incubation periods (48 hours), all *A baumannii* strains were capable of forming biofilms *in-vitro* at 44°C, which makes this pathogen as a succesful survivor in adverse conditions of higher temperatures in hospital environments.

Biofilm forming capacity varied among different clinical sources which showed respiratory isolates as strong biofilm former followed by pus then peritoneal fluid.

AST and MIC result discussion:

Majority of strains were associated with respiratory infections followed by blood, pus urine and peritoneal fluid. Antibiotics included for AST and MIC against bacterial isolates are based on their usage as conventional antimicrobial regimen prescribed by medical practitioners for treating *A. baumannii* infections. Therefore, levofloxacin, tetracycline, amikacin, ceftazidime, cefepime, imipenem, meropenem, piperacillin-tazobactam and cefoperazone-sulbactam were utilized in the experiment. AST and MIC results of antibiotics are interpreted according to CLSI guidelines 2015 through visible circular and elliptical zone of lysis around susceptible isolates against antibiotic disk and E strips.

Proportions of resistance among strains isolated from respiratory secretions are in the range between 68%-85%. While, highest proportions of sensitive *A. baumannii* isolates against all tested drugs are originated from pus and peritoneal fluid. *A. baumannii* susceptibility rates observed for different antibiotics were Levofloxacin (16.3%), Tetracycline (26.5%), Amikacin (28.5%), Meropenem (20.4%), Imipenem (14.2%), Ceftazidime (22.4%), Cefepime (16.3%), Piperacillin-tazobactam (18.3%), and Cefoperazone-sulbactam (20.4%).

Resistance rates were high for most antimicrobial agents with the exception of amikacin and to a lower extent to ceftazidime, meropenem and cefoperazone-sulbactam. Respiratory isolates were found most sensitive against amikacin & ceftazidime. Isolates from pus and peritoneal fluid showed highest level of resistance against all antibiotics (50-100%).

Proportions of sensitive isolates against all tested antibiotics were found between 15%-30%. Amikacin, ceftazidime, cefoperazone sulbactam showed highest sensitivity against *A. baumannii* isolates more than 80% resistance levels were observed against imipenem. Overall, antibiotic resistant strains from different sources expressed more ability to form biofilms as compared to antibiotic sensitive strains.

Motility assay:

While studying about motility features in biofilm-forming strains of *A. baumannii*, though very less is known about the motility for many of the strains, isolates demonstrating elevated motility might be due to in excess of expression of type IV pili-related genes as contrasted to non motile.

Otherwise non motile isolates that exhibit small or nix motility may be deficient of type IV biogenesis genes. 50% of peritoneal fluid isolated showed high motility, followed by respiratory sources i.e. 8.3%. Least or null motility was shown by pus derived strains may be due to lack of pili in their structures. These results contradict the initial facts about *A.baumannii* that it is a non-motile pathogen. These results also contribute towards the fact that motility of these organisms help them in colonization and biofilm formation on biotic and abiotic surfaces.

Antibiofilm activity of nanoparticles:

For 2 hours: sample 1 showed no effect on the biofilm capability of ATCC 1605 at any concentration.

Sample 2 showed decrease in biofilm formation at concentration of 1024 μ M, showing antibiofilm activity but showed comparatively high biofilm formation at a lower concentration of 256 μ M.

For 4 hours: sample 1 also showed decrease in biofilm formation at concentration of 1024 μ M, showing antibiofilm activity but showed high biofilm formation at a lower concentration of 0.5 μ M.

Sample 2 showed lesser biofilm production at concentration of 256 μ M as compared to biofilm formation at lesser concentration of 32 μ M, of the sample.

For 8 hours: both sample 1 and sample 2 had null or less effect on the biofilm forming capacity of ATCC 1605 and showed no positive results.

It can be concluded that by increasing the incubation time periods of the antibiofilm compounds their antibiofilm activity is reduced and deteriorated.

Chapter 6: References

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Chapter 7: Appendix

- 1x PBS [pH 7.2]: composed of 137 mM NaCl, 2.7 mM KCl, 10 mM Na₂HPO₄, and 2 mM KH₂PO₄.
- Modified luria broth: tryptone – 10 g/l; NaCl – 5 g/l; yeast extract – 5 g/l; agar- 0.4%.
- Ethanol:acetone :80ml of ethanol +20ml of acetone

- Luria broth: 20gm for 1000ml of distilled water.
- Mc farland reagent:

Mc Farland standard no.	0.5	1	2
1% barium chloride(ml)	0.05	0.1	0.2
1% sulphuric acid(ml)	9.95	9.9	9.8