



Characterization of Biofilm Production in Antibiotic Resistant *Klebsiella Pneumonia* Isolated From Different Clinical Samples in Iraqi Hospitals

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Abstract

Objective(s) *Klebsiella pneumonia* is a causative agent of chronic infections leading to increase morbidity and mortality; especially when performing biofilm (1). Therefore, biofilm formation has been linked to the survival of pathogenic bacteria in the hospital environment, leading to susceptible bacterial colonization which creates an important public health problem. Therefore, to characterize biofilm production in local Iraqi *K. pneumonia* isolated from different clinical samples in Iraqi hospitals using phenotypic biofilm assays including scanning electron microscopy. Material and Methods local isolates were collected from different Iraqi hospitals and re-diagnosed by compact Vitek 2 and genetically by using housekeeping gene (16s rRNA and 23s rRNA 639 bp). Phenotypic detection of biofilm formation among isolates was screened by using micro titer dish assay, twitching motility assay, and scanning electron microscopy (SEM). Results: It was found that 22/24 (91.67%) of isolates could form biofilm ($OD \geq 0.68$). Twitching motility test revealed that 20/24 (80.3%) of isolates could move on the M63 medium. Scanning electron microscopy showed 83.3% of *K. pneumonia* isolates made biofilm in different stages started with adhesion step and ending with a mushroom like architecture as highly magnification images showed on glass cover slips embedded statically in LB broth. It was interesting that *K. pneumonia* S2 demonstrated a dense mat of cells aggregates on cover slips generating different biofilm formation step. Conclusion *K. pneumonia* could produce biofilm, describing the ability to resist many kinds of antibiotics and rising a concern of disseminating chronic infection among hospitalized patients in many hospitals in Baghdad.

Keywords: Biofilm, Twitching motility, ELISA, Scanning electron microscopy.

Introduction

K. pneumoniae is a gram-negative, nonmetal, ubiquitous, facultative anaerobic and rod-shaped bacterium (1). It is also an opportunistic pathogen that can cause many clinical outcomes including UTIs, bacteremia, meningitis, skin and soft tissue infections, hospital and community-acquired pneumonia, ankylosing spondylitis, cholecystitis, osteomyelitis (2). *K. pneumoniae* has included among the six ESKAPE pathogens (3) responsible for roughly 70% of Gram-negative infections in hospital intensive care units (ICU) (4). It has emerged as a common cause of serious

epidemic and nosocomial infections in hospitals, resulting in high morbidity and mortality (5; 6). It is demonstrated that the mortality rate was 46/82 (56%) according to retrospective cohort study from tertiary care of bloodstream infection and 9/9 (100%) according to case-control study of nosocomial *K. pneumoniae* infections in Asia Pacific region (7). Biofilm producer *K. pneumoniae* could be resistant to wide range of antibiotics (8; 9). Biofilm is a microbially derived sessile community which is characterized by cells that are irreversibly attached to a substratum or interface with

each other, and embedded in a matrix of self-produced extracellular polymeric substances (10). Adaptation to surface attached growth within a biofilm is accompanied by significant changes in gene and protein expression, as well as metabolic activity which confers resistance to antimicrobial therapy and host mechanisms of clearance (10). Many pathogenic and nosocomial bacteria have been observed to predominantly exist as biofilms, in both natural environments and within infected tissues as polymicrobial communities (11).

Biofilm formation can be divided into distinct stages, from the initial attachment of bacteria to the surface to the formation of mature biofilm with a characteristic three-dimensional architecture (9). Bacterial functions such as motility, adhesion, transport, stress response, activation of metabolic pathways and extracellular matrix synthesis are required at each step of biofilm production (12).

In our study, we focus on the characterization of biofilm production in local Iraqi *K. pneumonia* isolated from clinical samples in Iraqi hospitals and tested for antibiotic resistant pattern in our previous work (13) using phenotypic biofilm assays including scanning electron microscopy (SEM).

Material and Methods

Collection and Diagnosis of Bacterial Isolates

Twenty four isolates of *K.pneumonia* were obtained from patients admitted to several teaching hospitals in Baghdad during a period between March 2015 to September 2015 from urinary tract infections (5 isolates), bacteraemia (7 isolates), wound swabs from Burn Unit (4 isolates), 2 isolates were collected from each sputum and ear, and pus, and finally 1 isolate collected from both of wounds infections and stool. Isolates were cultured on CHROM agar Orientation plates. Also, they were re-diagnosed by Vitek 2 compact automated system (Biomereux, USA), and the probability was 99% *Klebsiella pneumoniae ssp pneumoniae*.

Genotyping Detection for Isolates

Bacteria were diagnosed on the genetic level using amplified housekeeping gene (16s rRNA and 23s rRNA) designed using Geneious Software/ primer 3. The amplified size was 639 bp from the origin gene sequence F-3' TGTACACACCGCCCGTC-5'; R-3'GGTACTTAGATGTTTCAGTTC-5'. Total DNA was extracted using Genomic DNA Extraction Kit (Wizards, Promega, USA) following manufacture's protocol for Gram – negative bacteria from overnight bacterial growth.

The extract was used as a DNA template for the PCR process. The mixture of PCR is composed from 12.5 of Go Taq® Green Master Mix (2x), 5 µl template DNA, 1.5 µl primers (for each) final concentration (0.6 pmol/µl) and nuclease free water up to 25 µl (4.5 µl). Uniplex PCR products of the segment (16S rRNA and 23s rRNA) were visualized using 1% agarose gel electrophoresis.

Phenotypic Detection of Biofilm Production among Local Isolates

Microtiter Dish Assay

Biofilm formation in 96-well microtiter plates was assayed and quantified as previously described by O'Toole (14). All biofilm assays were performed using M63 minimal medium supplemented with glucose, MgSO₄, and CAA. Isolates were grown overnight in LB broth at 37°C, the 96-well plate(s) were prepared for the assay. Each isolate suspension was diluted (1:50) into an aliquot of the Biofilm media (M63) and mixed well by swirling and pipetting up and down. Standard strains *Pseudomonas* PA14 and *Pseudomonas* ΔPel A mutant were also undergone this step and used as positive and negative control, respectively.

The wells were inoculated (at least 4 wells per each isolate) of the 96-well plate (100 µl/well) from the isolate mixture using a multi-channel pipette. The 96-well plate was covered with a lid and incubated at (37°C) for up to 24 hours. After the incubation period, the wells were rinsed twice with distilled water to remove the unattached bacteria. Subsequently, 125 µl of Crystal violet stain (at 0.1% concentration) was added to each well and the control well. The plate was let sit to 10-15 min.

The plates were then rinsed twice with distilled water and dried on paper towels. Biofilm formed for each isolate was quantified by measuring their absorbance at 550nm using ELISA system. Quantify absorbance at 550 nm was read for the triplicate wells using 30% acetic acid in water as the blank.

Then the average and the standard deviation (SD) of absorbance values for each set of triplicate standards and triplicate samples were calculated. Since there is no universally recognized reference value used for evaluating biofilm formation capacity, in current study, isolates with OD₅₅₀ values greater than that of the negative control were considered positive for biofilm formation. Specifically, those with OD₅₅₀ values greater than that of the negative control, but less than that of the positive control were characterized as weak biofilm formers, while those with OD₅₅₀ values greater than that of positive control were considered strong biofilm formers.

Twitching Motility Assay

Each isolate was tested for its twitching motility character according to (15). Twitch motility plates consisted of M63 medium supplemented with MgSO₄, glucose (20%), CAA (20%), and solidified with 1.5% agar-agar. Isolates were stab inoculated with a toothpick through a thin (approximately 3 mm) LB agar layer supplemented with M63 to the bottom of the Petri dish and incubated at 37°C for 24 hours.

After incubation, the zone of twitching motility between the agar and Petri dish interface was visualized by staining with crystal violet. Results interpretation was dependent on categorizing these isolates into three groups according to positive (*Pseudomonas* PA14) and negative (*Pseudomonas* ΔPel A mutant) control. If the twitching zone diameter was <5 mm, the isolate is considered as twitching negative. A twitching zone diameter between 5 and 20 mm is considered as an intermediate while >20 mm of twitching zone was considered as a highly motile isolate.

Scanning Electron Microscopy SEM

A sterile cover slip was placed inside LB

broth tubes cultured with overnight bacteria. The tubes were incubated horizontally at 37°C/24 hrs, and then isolates were left for more than one month at room temperature in order to let a pellet of colonies to be formed. Isolates samples were prepared for testing by SEM according to the following steps provided by (16).

Each cover slip having biofilm formation appearance was removed from LB broth tube very carefully using sterilized forceps. Fixation step was done for the cover slip using a chemical fixative like glutaraldehyde (2.5%). The cover slips were incubated at room temperature for approximately 2 hours. The fixative agent was then removed and replaced immediately with distilled water to prevent sample dehydration. Cover slips were then coated with gold/ palladium (Au/Pd) Ratio (80:20) (57mm diameter x 0.1mm thick) using a sputter coater, plasma chambers with low discharge. Clear view of biofilm stages were conducted using a conventional scanning electron microscope (SEM) at a magnification from 5µm to 2mm.

Results

Identification of local Isolates

Results showed that all isolates exhibited on CHROM agar Orientation metallic blue colonies (figure 1-1), and also appeared to be *Klebsiella pneumoniae ssp. pneumoniae* at 99% level by Vitek 2 compact automated system. Results of genotypic detection of housekeeping gene 16SrRNA and 23SrRNA through isolates showed 100% positivity with amplified size of 639 bp.



Figure (1-1): *K. Pneumonias* isolates Showed Metallic Blue Colonies on CHROM Agar Orientation.

Micro titer Dish Assay

Biofilm formed by each isolate was quantified by measuring its absorbance at OD_{550nm} using ELISA system and

compared the results with positive (*Pseudomonas* P14) and negative control (*Pseudomonas* ΔPel A mutant) to be classified into either strong or weak biofilm producers (OD 0.35 and 0.05, respectively). Reading data demonstrated that 22/24 (91.7%) of *K. pneumoniae* isolates gave positive results as presented in table (1-1).

It was clear from the results that MDR *K. pneumoniae* U2 and S1 were not biofilm

producers compared to other MDR isolates that were weak biofilm producers; however, XDR *K. pneumoniae* S2 gave (OD≥0.68) when measuring its absorbance by ELISA system followed by XDR b4 and XDR b5 that gave (OD≥0.62 and 0.65, respectively); showing that XDR *K. pneumoniae* isolates gave the highest rate of biofilm production using this method (figure 1-2).

Table (1-1): Results of phenotypic detection method of biofilm among 24 *K. pneumoniae* isolates.

No. of isolates	Phenotypic detection of biofilm formation among isolates		
	Microtiter Dish assay readings by ELISA at 550nm average OD± SD	Biofilm formation pattern	Twitching motility assay mm
Positive control (<i>Pseudomonas</i> PA14)	0.4±0.001	Strong	3
Negative control (<i>Pseudomonas</i> ΔPel A mutant)	0.05±0.02	None producer	None producer
MDR <i>K. pne.b1</i>	0.33±0.02	weak	6
MDR <i>K. pne.b2</i>	0.31±0.1	weak	9
MDR <i>K. pne.b3</i>	0.45±0.001	strong	6
XDR <i>K. pne.b4</i>	0.62±0.01	strong	25
XDR <i>K. pne.b5</i>	0.65±0.01	strong	30
XDR <i>K. pne.b6</i>	0.53±0.01	strong	22
MDR <i>K. pne.b7</i>	0.32±0.004	weak	3
MDR <i>K. pne.U1</i>	0.14±0.01	weak	0
MDR <i>K. pne.U2</i>	0.01±0.02	none	13
XDR <i>K. pne.U3</i>	0.44±0.02	strong	10
MDR <i>K. pne.U4</i>	0.33±0.01	weak	7
MDR <i>K. pne.U5</i>	0.32±0.01	weak	3
MDR <i>K. pne.S1</i>	0.02±0.01	None	0
XDR <i>K. pne.S2</i>	0.68±0.01	Strong	35
MDR <i>K. pne.W1</i>	0.44±0.02	Strong	23
XDR <i>K. pne.ES1</i>	0.46±0.02	strong	11
XDR <i>K. pne.ES2</i>	0.56±0.01	strong	25
MDR <i>K. pne.Bu1</i>	0.27±0.01	weak	10
XDR <i>K. pne.Bu2</i>	0.56±0.02	strong	20
MDR <i>K. pne.Bu3</i>	0.59±0.01	strong	15
XDR <i>K. pne.Bu4</i>	0.66±0.004	strong	28
XDR <i>K. pne.P1</i>	0.56±0.01	Strong	23
MDR <i>K. pne.P2</i>	0.26±0.01	Weak	7
MDR <i>K. pne. St1</i>	0.48±0.03	Strong	24

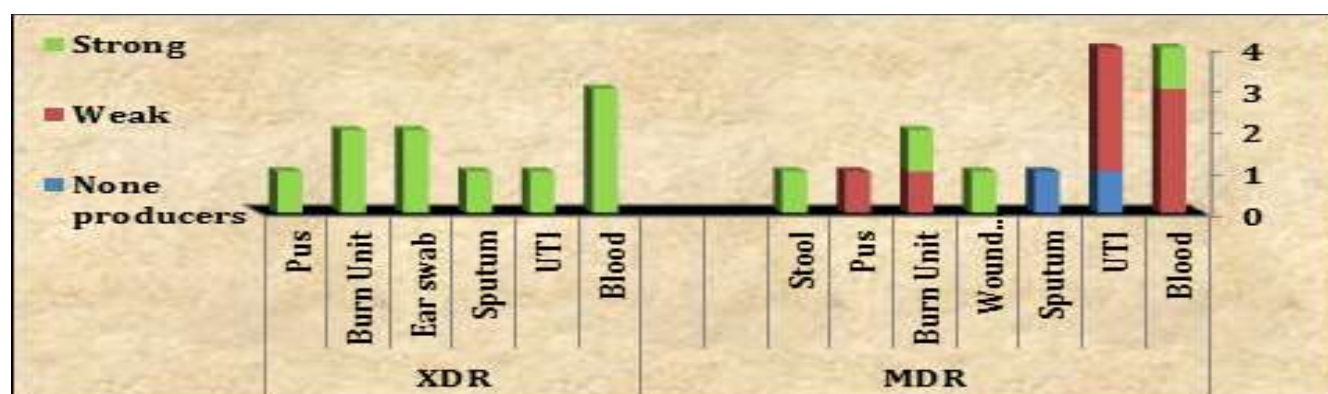


Figure (1-2): Classifying the MDR and XDR *K. pneumoniae* isolates into none-, weak, or strong biofilm producers. Results obtained from microtiter dish assay in current study

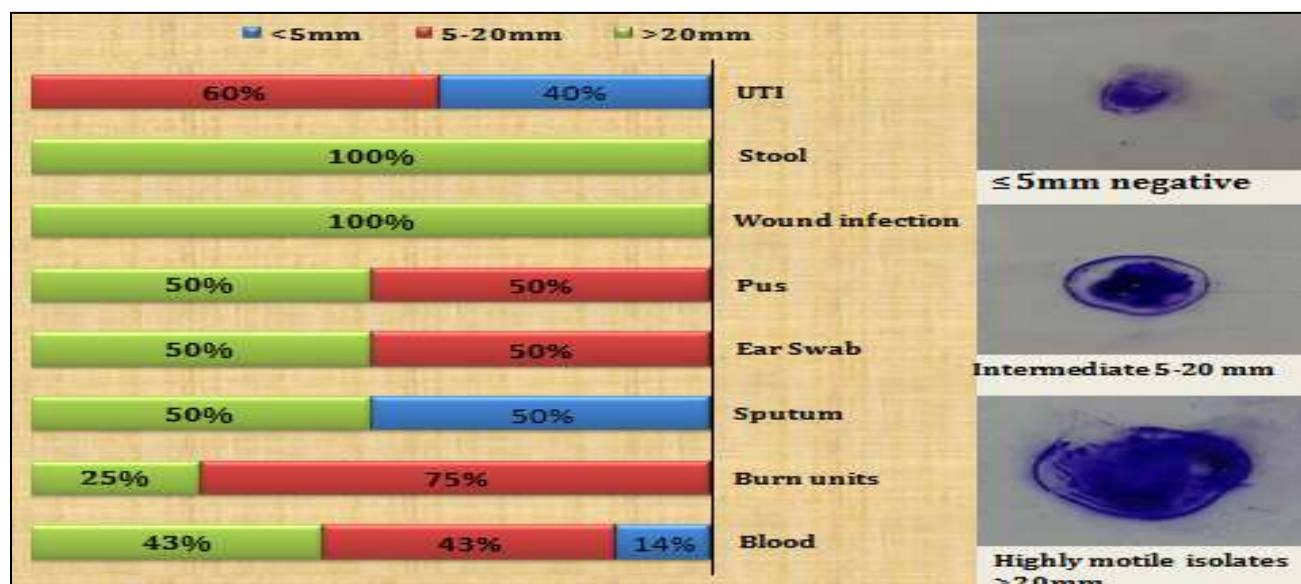


Figure (1-3): Twitching motility profile presented as percentages (left) among 24 *K. pneumoniae* isolates collected from different clinical samples and migration zone on M63 agar plates (right) for *K. pneumoniae* U1 (3mm/negative), ES1 (11mm/intermediate), and S2 (35mm/high motility).

Twitching Motility Pattern

The twenty four *K. pneumoniae* isolates were also tested for their twitching motility pattern on M63 agar plates. Results demonstrated that there was 20/24 (80.3%) of *K. pneumoniae* isolates exhibited variable twitching motility patterns after incubation period (Table 1-1). Figure (1-3) represent the variable percentages of twitching motility profile among different clinical sources. XDR *K. pneumoniae* S2 found to be the highly motile isolate that exhibited 35mm in diameter immigration zone on M63 agar plates, followed by XDR Bu4 and XDR b5 which showed 30 and 28mm immigration zones, respectively.

Scanning Electron Microscopy (SEM)

The twenty four antibiotic resistant *K. pneumoniae* isolates were also undergone SEM analysis to have an understanding view of biofilm formation among those isolates related to different clinical samples obtained locally. SEM analysis showed that 20/24 *K. pneumoniae* isolates formed biofilm with different stages on glass liquid interface.

It was interesting that each biofilm producing *K. pneumoniae* isolate demonstrated a dense mat of cells aggregates

after the incubation period on cover slips (figure 1-4). This tight cell aggregate could be due to factors involved in aggregative adhesions (type 1 or type 3 fimbriae), luxuriant capsules that all *K. pneumoniae* isolates exhibited earlier, and the lipopolysaccharides (LPS) which allow bacteria to grow as a biofilm. Highly magnification images presented in figure (1-4 No.1) showed condense monolayer formed by XDR *K. pneumoniae* S2 which generated a large amount of extracellular components that interact with organic and inorganic molecules in the LB medium to create the glycocalyx, consequently forming microcolonies as shown in No.2 in the same figure.

Mature biofilm architecture was also identified for this bacterium and found to be heterogeneous, containing microcolonies of bacterial cells embedded in an exopolysaccharide (EPS) matrix and separated from other microcolonies by water channels that allow easy distribution of nutrients and oxygen (figure 1-4No.3). The property of highly antimicrobial resistant *K. pneumoniae* attachment and high rate of biofilm formation on glass showed by SEM placed them among the infectious bacterial pathogens that could be most challenging to control

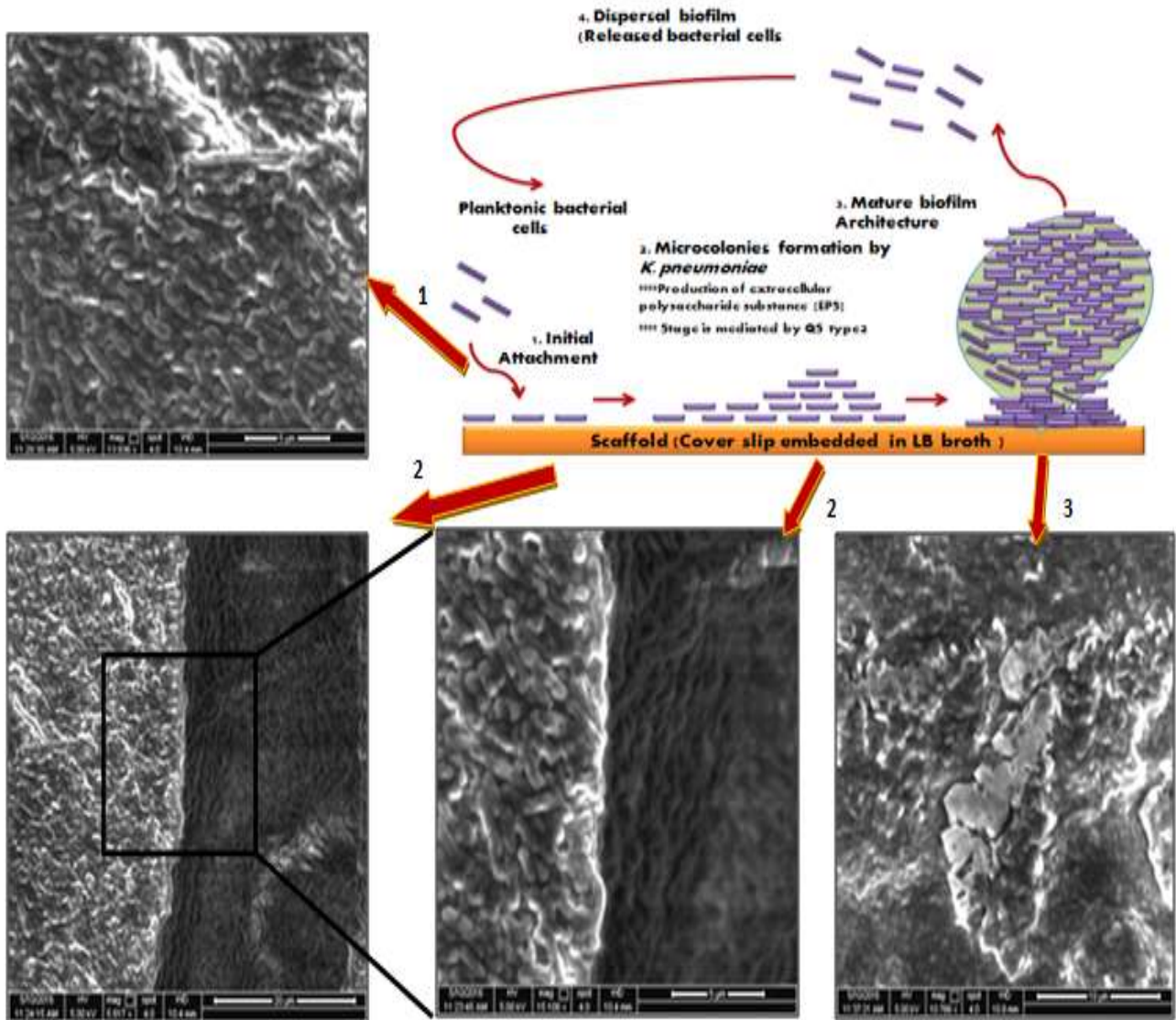


Figure (1-4): SEM images of different biofilm formation stages of *K.pneumoniae* S20 on glass cover slip; 1) Adherent stage and attachment of cells tightly (x5µm), 2) (x5µm and x20µm) of irreversible stage presented in microcolonies; 3) Mature stage of biofilm at (x10µm) magnification where cells aggregated in 3D shape that had channels for water and nutrition

Discussions

All *K. pneumoniae* isolated from different clinical samples gave the right color for Enteriobacteriaceae family on CHROM agar Orientation. Same observation was detected by Ajao *et al.*(17) and Manickam *et al.*(18) who both further revealed that degraded chromogens would allow easy identification of mixed growth and provides higher detection rates. Also, Vitek2 Compact automated system was used in current study since the importance of accurate identification of specific species becomes paramount in most modern microbiology laboratories and relies heavily on automated systems (12); especially this automated system which provides rapid and accurate

identification to the species level (19). Genotypic detection showed 100% positivity among isolates with amplified size of 639 bp, supporting what had been explained for *K. pneumoniae* identification by phenotypic methods above. The identification to the species level was known to be difficult due to similarity of biochemical profiles among species for the same genus, such as *K. planticola* or *K. oxytoca* and *K. pneumoniae* subsp.

pneumoniae that cannot be differentiated by phenotypic tests (20). Consequently, conventional PCR has been used to amplify the highly conserved rRNA genes (16S, 23S, and 5S) which are ideal candidates for

bacterial identification and evolutionary studies (21). As reported from sequence analysis studies, 16S and 23S rRNA could be used to identify *K. pneumoniae* subsp. *pneumoniae* to the species level (22). Besides, Dong *et al.*(23) postulated the importance of 16S-23S internal transcribed spacer to detect *K. pneumoniae* at species level.

Results of microtiter plates showed the capability of different *K. pneumoniae* isolated from clinical source to produce either strong or weak biofilm. Strongest biofilm formed by *K. pneumoniae* isolated from sputum followed by isolate from bacteriemia. Results agreed with Langstraat *et al.*(24) and Jagnow and Clegg (25). Martino *et al.*(26) also demonstrated that only 48% of *K. pneumoniae* isolates were moderate biofilm forming strains on polystyrene dishes.

However, Maldonado *et al.* (27) postulated the capability of different *Klebsiella* isolates from clinical source to produce biofilm. They found that *Klebsiella pneumoniae* subsp. *pneumoniae* isolated from UTIs was able to form biofilm (OD > 0.5) at 550nm. Olewi and Abid (28) showed that 8 *K. pneumoniae* isolated from drinking water in different areas of Baghdad can strongly produce biofilm with mean of (OD > 0.7).

Results of twitching motility assay showed that MDR *K. pneumoniae* isolate U2 and S1 could not exhibit any motility pattern along with MDR *K. pneumoniae* b7 and U1 which showed motility for less than 5mm, and this would agree with their results in microtiter dish assay; concluding that they were not biofilm producers.

Few studies in literature have been addressing the twitching motility of *K. pneumoniae*; however, Liaquat *et al.* (29) observed twitching motility in order to determine any effect in the biofilm forming capability of *K. pneumoniae* isolates. They further revealed that all the isolates tested were efficient biofilm-formers and had this motility pattern.

Highly magnification images achieved by Scanning electron microscopy showed that local *K. pneumoniae* isolates were capable to produce strong biofilm with different stages;

especially the interesting tight mate of extensively *K. pneumoniae* isolated from sputum specimen attached to each other and to glass substratum. Results were supported by Balestrino *et al.* (30) and Pour *et al.* (31).

It was further reported that biofilm characteristic of *K. pneumoniae* isolates causing pneumoniae were most studied due to enhance the ability of this bacterium to transfer resistance markers to other clinical strains in mixed infection (31). Abd El-Baky (32) postulated that 22% of *Klebsiella* spp. was the most prevalent microorganism examined for biofilm formation using SEM. It was also reported that there were denser and high cell aggregates on cover slips leading to microcolonies formation that separated by water channels (33).

Both Bellifaet *et al.* (34) and Jamal *et al.*(35) demonstrated that most biofilm forming *K. pneumoniae* strains were strongly adhere to glass slides and were at least 10 times more antibiotic resistant than their planktonic counterparts. They referred the high adherence with the presence of type 3 fimbriae (36; 37).

Conclusion

Phenotypic detection and scanning electron microscopy (SEM) of local *K. pneumoniae* isolated from different clinical samples among hospitals in Baghdad indicated big concern of disseminating chronic infection among hospitalized patients; especially biofilm producer *K. pneumoniae* that could resist wide range of antibiotics by various methods.

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Abbreviation

MDR: Multidrug resistant

XDR: Extremely drug resistant

PDR: Pan drug resistant**SEM:** Scanning electron microscopy**QS:** Quorum sensing**References**

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