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## Detection of Biofilm Formation Among the Clinical Isolates of *Klebsiella pneumoniae*: Phenotypic and Genotypic Methods

Fatimah E. Alquraishi<sup>1\*</sup>, Zainab N. AL-Saadi<sup>1</sup> and Jalal A.T. Al-Azzawi<sup>2</sup>

<sup>1</sup>Department of Biology, College of Science, University of Wasit, Iraq.

<sup>2</sup>Ministry of Health, Wasit Health Directorate, Wasit Governorate, Iraq.

\*Correspondence: fatimaenad6@gmail.com; Tel.: (+09647726878654)

### Abstract

Infections caused by biofilm-embedded pathogens decrease the efficacy of traditional treatments and increase antibiotic tolerance. Most of the human bacterial infections are biofilm-associated. Therefore, this study aimed to detect the biofilm formation among the clinical isolates of *Klebsiella pneumoniae* collected from different hospitals in Wasit province-Iraq by phenotypic and genotypic methods. 525 clinical samples were used to isolate 77 *K. pneumoniae* strains from clinical specimens for five months. They were identified by microbiological method as *K. pneumoniae*. The microtiter plate method is used to detect the biofilm formation. Results showed that out of 77 *K. pneumoniae* isolates, 76 (98.7%) isolates were biofilm producers with three different categories; 12 (15.6%) were weak-biofilm producers, while other isolates 63 (81.8%) and 1 (1.3%) were moderate and vigorous producers, respectively. However, 1 (1.3%) isolates were identified as non-biofilm producers. Amplification of genes by multiplex PCR technique was done for 77 isolates of *K. pneumoniae* to detect biofilm production genes, *mrkD* and *FimH*. Results showed that out of 77 isolates, there were 74 isolates (94.8%) positive to *mrkD* and 33 isolates (42.8%) to *fimH*.

**Keywords:** *K. pneumoniae*; Microtiter plate method; *mrkD*; *fimH*; Iraq.

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### Introduction

*Klebsiella pneumoniae*, a member of the family *Enterobacteriaceae*, is a part of the flora and is isolated as the causative agent in complicated infections. It is a microorganism that causes severe diseases such as pneumonia, septicemia, bacteremia, wound infections and purulent abscesses at different sites in humans. The bacterium is widely distributed in healthy people's urinary, respiratory, and gastrointestinal tracts. Most *K. pneumoniae* are hospital-associated with a high fatality rate if incorrectly treated<sup>1</sup>. *Klebsiella* spp. is among the five gram-negative pathogens most usually encountered in hospital-acquired infections.<sup>2</sup> *Klebsiella* was found in two general habitats: the first habitat is the mucosal surface of mammals such as humans, swine or horses, which they colonize, and the other habitat is the environment, where they are found in surface water, soil and on plants<sup>3</sup>. *Klebsiella pneumoniae* can produce several virulence factors necessary for the infection's colonization, adherence, invasion and progress. These include lipopolysaccharide presence, biofilm production, capsule production, adherence factors and siderophore activity. In addition, further virulence factors such as

hemolysins, protein tyrosine kinase, heat-stable endotoxins and heatlabile exotoxin<sup>4,5,6</sup>.

Filament formation is a typical response in which bacteria replicate but incompletely divide, leading to long, slender chains. Many bacteria have existed in biofilm for prolongation. Biofilms preserve persistent bacteria from antibiotic effects, leading to the emergence of persistent bacterial cells. It should be noted that persistent pathogens lead to chronic diseases, resulting in the overuse of antibiotics and reduced antibiotic efficacy<sup>7</sup>.

## Materials and Methods

### Specimens collection

This study was carried out in Al-Haj Jalal Hospital in Wasit Province-Iraq. 525 specimens from patients clinically diagnosed with UTI from 1<sup>st</sup> October to 30<sup>th</sup> February 2021. Of these patients, 442 were females (aged 25 days to 80 years), and 83 were males (aged 1 month to 80 years). All urine samples were collected in sterile screw-capped test tubes.

### Isolation and identification

Clinical samples were cultured onto MacConkey and Blood agar plates for 18-24 h at 37°C. Morphologic characteristics tested all lactose-fermenting isolates according to MacFaddin<sup>8</sup>. The collected isolates were identified biochemically according to Forbes et al.<sup>9</sup> and methods described by MacFaddin<sup>8</sup>. Confirmation of *K. pneumoniae* was conducted using the API20E system. The test was done according to the manufacturer's instructions (BioMeriux, France). Bacterial isolates were stored in BHI broth containing 20% glycerol at -20°C<sup>10</sup>.

### Detection of biofilm formation

#### Phenotypic method

Biofilm formation test was done by microtiter plate method as described by Al-Timimi<sup>11</sup>. Bacterial isolates were cultured on brain heart infusion agar and incubated at 37°C for 24 h. then, a few (3-5) colonies, suspended in 5 ml of normal saline in test tubes, were mixed by vortex. Twenty microliters of bacterial suspension overnight culture were used to inoculate 96 wells of a bottomed microtiter plate containing 180 µl of brain-heart infusion broth with 2% sucrose. Control wells contained 200 µl of BHI broth with 2% sucrose; triplicate was done for each isolate. The microtiter plate was covered with a lid during incubation at 37°C for 24 h. After incubation, the content of each well was removed, and the wells were carefully washed three times with PBS (pH 7.2) and then left to dry. The plates were dried at room temperature for 15 minutes. Crystal violet (1%) was added to the wells for 15 minutes. After removing the crystal violet solution, wells were washed three times with PBS (pH 7.2) to remove the unbounded dye and allowed to dry at room temperature. Dye bound to the adherent cells was disbanded with 200 µl ethanol. The absorbance of each well was measured at 630 nm using an ELISA reader. The control well's optical density (OD) value was deducted from the entire test, and each assay was performed in triplicate. The adherence capabilities of the bacterial isolates were classified into four categories; above, the mean optical density of the negative control (contained broth only) was considered as the CUT-OFF, and isolates were classified according to Mathur et al.<sup>12</sup> as follows:

#### OD value Biofilm ranking

$OD \leq OD_c$ .....	None
$OD_c < OD \leq 2 \times OD_c$ .....	Weakly
$2 \times OD_c < OD \leq 4 \times OD_c$ .....	Moderate
$4 \times OD_c < OD$ .....	Strong

### Genotypic method

DNA was extracted from 77 *K. pneumonia* clinical isolates using a commercial purification system Easy Pure<sup>®</sup> Bacteria Genomic DNA Kit. The extraction of genomic DNA was performed according to the company's manufacturing. Aseptically prepared the PCR reaction mix using Taq Ready master mix Kit according to the manufacturer's instructions for a final reaction volume of 50 µl with 7 µl of DNA extract. Multiplex PCR of each primer was performed with Taq GreenMaster Mix PCR Kit. Reaction mix 50 µl consisted of 25 µl of 1X PCR Master Mix, 7.5 µl of biofilm formation genes (3.75 µl from *mrkD*, and 3.75 µl from *fimH*) of each primer, and 7 µl (10–100 ng) of template DNA and then complete the volume into 50 µl by nuclease-free water. DNA amplification was carried out with the following thermal cycling: an initial denaturation of DNA at 95°C for 15 min. was followed by 35 cycles of amplification (95°C for 40 sec, 52°C for 30 sec, and 72°C for 45sec), ending with a final extension at 72°C for 5 min, and soak at 4°C for 5 min. The sequence of oligonucleotide primers was used to detect *fimH* [F (5'-CGGAAACGATCACCGACTAC-3') and R (5'-CACGTCGTTATTGGCGTAGA-3')] and *mrkD* [F (5'-CCACCAACTATTCCCTCGAA-3') and R (5'-GGCCGACGGTGTATTCTTA-3')] at a product size of 489bp and 317bp, respectively.

Electrophoresis results were identified using a UV-Transilluminator system. The DNA bands were measured according to the ladder DNA. The positive results were distinguished when there was a DNA band equal to the target product size and then photographed using a camera

### Statistical analysis

This study's data results were analyzed using Graph Pad Prism 8 software and Microsoft Excel 2013 for each biological replicate. The probability level at P values below  $\leq 0.05$  was used to identify a significant difference<sup>13,14,15</sup>.

## Results

### Culture

Of 525 patients clinically diagnosed with UTI, only 233 (44.38%) were positive for bacterial culture (only one specimen was selected per patient).

### Morphological characteristics

For identification of *K. pneumonia* isolates based on morphological characteristics of the colonies on MacConkey agar and blood agar, the isolates appeared as large, mucoid and pink on MacConkey agar due to lactose fermenting. In contrast, they appeared white, large, and mucoid colonies on blood agar without hemolysis.

### Biochemical tests

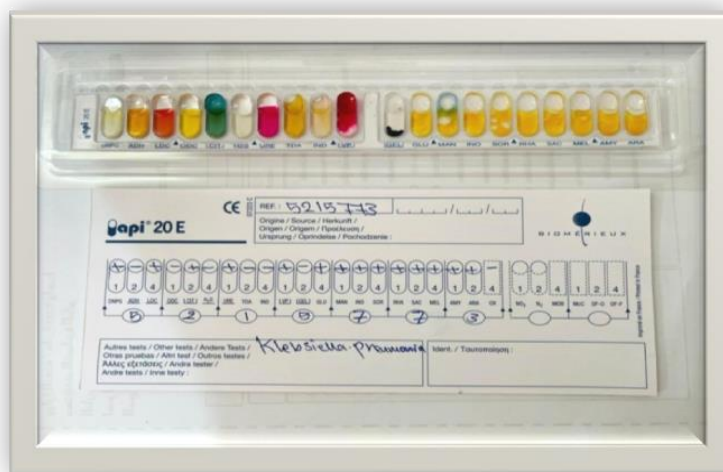
The biochemical tests were used to identify bacterial isolates (Table 1). It showed that all isolates of *K. pneumonia* revealed a positive catalase test indicated by bubbles formation of O<sub>2</sub>. While the results of IMViC differentiate them from other lactose fermenter genera showed negative results for indole. A positive result for the citrate utilization test utilization of citrate is an important physiological test. *Klebsiella* showed positive reactions to citrate. (Figure 1).

No.	Test	Result
1	Vogesproskauer	+
2	Indol test	-
3	Citrate utilization test	+
4	Methyl red	-

5	TSI	A/A/G+/H2S-
6	Oxidase	-
7	Catalase	+
8	Urease test	+

Positive result (+) ; Negative result (-); A (Acid); G+ (Gas production); H<sub>2</sub>S (No black sediment).

**Table 1. Biochemical tests used for confirming the identification of *K. pneumoniae* isolates.**



**Figure 1. API 20E system for *K. pneumoniae* characterization.**

Of 233 patients with positive bacterial culture, isolated *E. coli* from 79 (33.05%), followed by *K. pneumoniae* 77 (32.22%), *Staphylococcus aureus* 65 (27.2%), *Citrobacter* 6 (2.51%), *Proteus* spp. 5 (2.09%), *Enterobacter* 3 (1.26%), and *P. aeruginosa* 3 (1.26%), and *Streptococcus Pneumoniae* 1 (0.42%) (Table 2). The results of this study agreed with other local studies, such as Essa et al. [19] performed a study in Baghdad and Arbil on pregnant women, isolated *E.coli* from 49.1% followed by *Acinetobacter baumannii* (21.3%), *K. pneumoniae* (13%), *P. mirabilis* (11.1%), and *P. aeruginosa* (5.6%).

No.	Test	Result [No. (%)]
1	<i>Escherichia coli</i>	79 (33.05%)
2	<i>Klebsiella pneumoniae</i>	77 (32.22%)
3	<i>Staphylococcus aureus</i>	65 (27.2%)
4	<i>Citrobacter</i>	6 (2.51%)
5	<i>Proteus</i>	5 (2.09%)
6	<i>Enterobacter</i>	3 (1.26%)
7	<i>Pseudomonas aeruginosa</i>	3 (1.26%)
8	<i>Streptococcus Pneumoniae</i>	1 (0.42%)
9	Total	239

**Table 2. Positive bacterial culture distribution among 233 patients with UTI.**

The highest frequency of UTI was among young women (15-40 years), as shown in Table (4).

Gender	Age group of patients	Result [No. (%)]
Female	1month-14 years	15 (19.5%)
	15-40 years	(50.6%) 39
	>40 years	6 (7.7%)
<b>Total</b>		60 (77.8%)
Male	1month-14 years	8(10.4%)
	15-40 years	7 (9.1%)

	>40 years	2 (2.6%)
<b>Total</b>		17 (22.1%)

**Table 3. Gender and age distribution of *K. pneumoniae* isolates from patients with UTI.**

### *Biofilm formation*

#### Phenotypic method

*Klebsiella pneumoniae* forms biofilms on surfaces, whether biotic or abiotic, such as catheters and other medical devices, contributing to antibiotic resistance. Biofilms facilitate persistence, leading to persistent UTIs, which can lead to stone formation. Virulence factors like adhesions. From this point, the microtiter plate method tested the ability of *K. pneumoniae* isolates to produce biofilms (Table 4, Figure 2).

No. of isolate	Biofilm pattern	No. of isolate	Biofilm pattern
1	Moderate	40	Weak
2	Moderate	41	Moderate
3	Moderate	42	Moderate
4	Moderate	43	Weak
5	Moderate	44	Moderate
6	Weak	45	Moderate
7	Moderate	46	Non
8	Moderate	47	Moderate
9	Moderate	48	Weak
10	Moderate	49	Moderate
11	Moderate	50	Moderate
12	Moderate	51	Moderate
13	Moderate	52	Moderate
14	Weak	53	Moderate
15	Moderate	54	Moderate
16	Moderate	55	Moderate
17	Moderate	56	Moderate
18	Moderate	57	Moderate
19	Moderate	58	Moderate
20	Weak	59	Moderate
21	Moderate	60	Moderate
22	Moderate	61	Strong
23	Moderate	62	Moderate
24	Moderate	63	Moderate
25	Moderate	64	Moderate
26	Moderate	65	Moderate
27	Moderate	66	Weak
28	Moderate	67	Moderate
29	Moderate	68	Moderate
30	Moderate	69	Moderate
31	Moderate	70	Moderate
32	Moderate	71	Moderate
33	Moderate	72	Moderate
34	Weak	73	Moderate
35	Weak	74	Moderate
36	Weak	75	Moderate
37	Moderate	76	Moderate
38	Weak	77	Moderate
39	Weak	-	-

**Table 4. Biofilm pattern among isolates of *K. pneumoniae*.**

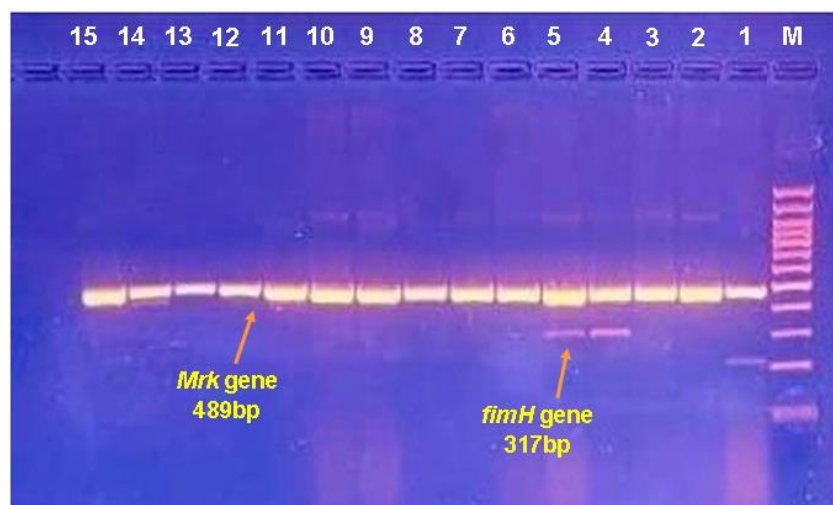


**Figure 2.** Microtiter plate test of biofilm-positive *Klebsiella pneumoniae*.

All 77 *K. pneumoniae* isolates were detected for biofilm formation by microtiter plate assay. Results showed that out of 77 *K. pneumoniae* isolates, 76 (98.7%) isolates were biofilm producers with three different categories: 12 (15.6%) were weak-biofilm producers, while other isolates 63 (81.8%) and 1 (1.3%) were moderate and strong producers, respectively. However, 1 (1.3%) isolates were identified as non- biofilm producers.

#### Genotypic method

*Klebsiella pneumoniae* isolates were genotyped, typically using the multiplex-PCR. Amplification of genes by multiplex PCR technique was done for 77 isolates of *K. pneumoniae* to detect biofilm production genes: *mrkD* and *FimH*. Results showed that out of 77 isolates, there were 74 isolates (94.8%) positive to *mrkD* 33 isolates (42.8%) to *FimH* (Figure 3).



**Figure 3.** The agarose gel electrophoresis image showed the multiplex PCR product of two genes (*fimH* and *mrk*) of *Klebsiella pneumoniae* isolates at 317 and 489 bp PCR product size, respectively. The Lane (M): DNA marker (100-1500bp).



## Discussion

This result was consistent with other Iraqi researchers. A study by Alsamarai and Ali<sup>12</sup> in Tikrit reported that 234 out of 563 (41.6%) gave positive cultures. In addition, Al-Jemely<sup>16</sup> in Baquba 135 urine samples, only 110 Samples gave bacterial growth at 81.4 %. Also, on Day<sup>17</sup> in Wasit, 278 out of 774 specimens (35.9%) were positive for bacterial culture. The findings of morphological characteristics were similar to those reported by another study<sup>16</sup>. The biochemical tests' results were identical to those of others<sup>8,17,18,19</sup>. Among UTIs caused by *K. pneumoniae*, females had a higher frequency than males. The results agreed with other studies. Bachay (2018) found that the highest frequency of UTI (51.7%) occurred in women aged 15-40 years in Iraq<sup>17</sup>. Iranpour et al. (2015) clarified that 50% of patients with UTI aged 15-45 years in Iran<sup>20</sup>. Bachay performed a study in Wasiton women, isolated *E. coli* from 67.20%, followed by *K. pneumoniae* (15.90%), *Proteus* spp. (6.70%), *P. aeruginosa* (6.70%) and *Enterobacter* (3.36%)<sup>17</sup>.

Concerning the phenotypic method, our results were identical to that detected by others<sup>21-25</sup>. The emergence of weak and moderate biofilm phenotypes among Gram-negative bacteria was in agreement with some local studies. Al-Rubyaie<sup>26</sup> indicated that all clinical *K. pneumoniae* isolates were able to produce biofilm 100%. Al-Timimi<sup>11</sup> observed that out of 50 *K. pneumoniae*, 40 isolates were biofilm producers with different categories, including 23 (46%), 14 (28%) and 3 (6%), which were weak, moderate and robust biofilm production, respectively. Also, Al-Husseini<sup>27</sup> observed that out of 100 *P. aeruginosa*, 85 isolates were biofilm producers with different categories, including 44% weak, 35% moderate and 6% strong biofilm production.

The genotypic result disagreed with Iraqi research, such as Al-Musawi<sup>28</sup>, in Baghdad, which reported *mrkD* in (82.85%) and a study done by Mirzaie and Ranjbar<sup>29</sup> in Iran, who reported *mrkD* in 88%. Our results agreed with Ferreira et al., 30 reported *mrkD* in 96% of Brazil. The result of *FimH* genes disagreed with local studies such as Al-Aajem et al.<sup>31</sup> in Diyala reported *fimH* genes in rates (86.66%), and Aljanaby and Alhasani<sup>4</sup> indicated *fimH* genes in rates 100%. Another study, such as Alsanie<sup>32</sup> in Saudi Arabia, indicated *fimH* genes in rates (69.5%). In India, Remya et al.<sup>33-34</sup> reported *fimH* genes in rates (84%)<sup>35</sup>.

## Conclusions

This study aimed to detect the biofilm formation among the clinical isolates of *Klebsiella pneumoniae* collected from different hospitals in Wasit province-Iraq by phenotypic and genotypic methods. The results support previous studies such as Essa et al.<sup>[19]</sup>. This section is mandatory but can be added to the manuscript if the discussion is unusually long or complex. All samples showed a positive result for urease production due to the ability to produce the enzyme urease; these enzymes converted the color from yellow to pink. All isolates were confirmed by using the API 20E system. Among UTIs caused by *K. pneumoniae*, females had a higher frequency than males regarding age and gender.

**Author Contributions:** for Conceptualization, FEA. and ZNA.; methodology, FEA. and JATA; software, ZNA.; validation, FEA., ZNA. and JATA.; formal analysis, FEA.; investigation, FEA. and ZNA.; resources, FEA.; data curation, FEA.; writing-original draft preparation, FEA.; writing review and editing, ZNA.; visualization, ZNA.; supervision, ZNA.; project administration, FEA. All authors have read and agreed to the published version of the manuscript.

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**Institutional Review Board Statement:** The study was conducted according to the guidelines of the Declaration of Helsinki and approved by the Ethics of the Scientific Committee of the Department of Biology in the College of Science (University of Wasit, Wasit, Iraq).

**Informed Consent Statement:** Informed consent was obtained from all subjects involved in the study.

**Conflicts of Interest:** The authors declare no conflict of interest.

## References

1. Martin RM, Bachman MA. Colonization, Infection, and the Accessory Genome of *Klebsiella pneumoniae*. *Frontiers Cell Infect Microbiol* **2018**; (8)4: 1-15.
2. Pokra M, Sharma DK, Mehta P, Verma HR, Pundir S, Rana J, Kulmi D. Its Alarming, *Klebsiella spp.* Towards Multidrug Resistance. *Int J Curr Microbiol App Sci* **2016**; 5(6): 150-60.
3. Goldman E, Lorrence HG. Practical Handbook of Microbiology. 2<sup>nd</sup> ed. USA. **2009**; pp: 21.
4. Aljanaby AAJ, Alhasani AHA. Virulence factors and antibiotic susceptibility patterns of multidrug resistance *Klebsiella pneumonia* isolated from different clinical infections. *Afr J Microbiol Res* **2016**; 10(22): 829-43.
5. Farhan Haneen Saad, Al-Saadi Zainab N, Ali Jassim Fatehi. Genotypic, Phenotypic identification and antibiogram assay of MRSA isolated from healthy individuals in Wassit province. *Indian J Forensic Med Toxicol* **2020**; 14 (4): 2043-2049.
6. Gharban HA. Cumulative Effect of Subclinical Mastitis on Immunological and Biochemical Parameters in Cow Milk. *Arch Razi Instit* **2021**; 76(6), 1599-1608.
7. Al-Saadi, Z. N. Bactericidal Activity of New Silver Nanoparticles Biosynthesis From Methicillin-Resistant *Staphylococcus aureus* Against Clinical MRSA and MSSA Isolates. *Biochem Cell Arch* **2020**; 20 (2): 5773-5780.
8. MacFaddin JF. Biochemical Tests for Identification of Medical Bacteria. 3<sup>rd</sup> ed, Lippincott Williams and Wilkins, London. **2000**; pp: 49.
9. Forbes BA, Sahm DF, Weissfeld AS. Bailey and Scott's Diagnostic Microbiology. 11<sup>th</sup> edition, Mosby, St. Louis, Mo, USA. **2002**; pp:156.
10. Al-Saadi ZN, Kasim FH, Al-Fatah A. Isolation and identification of bacteria causing respiratory tract infections in Tikrit city and determining its antibiotic resistance. *Tikrit J Pure Sci* **2005**; 10 (1): 16-20.
11. Al-Timimi SN. Persistence and Filaments Formation in *Klebsiella Pneumonia* Clinical Isolates. M.Sc. Thesis. Mustansiriyah University. **2021**; pp: 51.
12. Alsamarai AM, Ali S. Urinary tract infection in female in Kirkuk, Iraq: risk factors. *World J Pharm Pharm Sci* **2016**; 5 (6): 180-195.
13. Gharban HAJ, Al-Shaeli SJJ. Clinical and serum biochemical evaluation of goats with hypomagnesemia. *Biochem Cell Arch* **2021**; 21 (1): 587-592.
14. Mohammad HA, Ajaj EA, Gharban HA. The first study on confirmation and risk factors of acute and chronic canine distemper in stray dogs in Wasit Province, Iraq, using enzyme-linked immunosorbent assay and reverse transcription-polymerase chain reaction. *Vet World* **2022**; 15 (4): 968-974.
15. Razooqi MA, Gharban HA, Al-Kaabi MA. Molecular and Seroprevalence of Toxoplasmosis in Goats' Blood and Milk in Iraq. *Arch Razi Instit* **2022** 77(5): 1749-1755.
16. Al-Jemely HTN. A comparative study of the bacterial virulence factors isolated from urinary tract infections for cardiac care unit patients. M.Sc. Thesis. Diyala University. **2017**; pp: 118.
17. Bachay Z. Molecular Comparison of Adhesins and Phylogroups of Uropathogenic *Escherichia coli* Isolates From Patients with First Time and Recurrent Urinary Tract Infection. MSc Thesis. Wasit University. **2018**; pp: 43.
18. Abedalhammed, H. S., Naser, A. S., Al-Maathedy, M. H., Mohammed, Th. T., Jaber, B. T. & Al-Asha'ab, M. H. The effect of vitamin e as an antioxidant with different levels of dried tomato pomace supplementation on diets of common carp (*Cyprinus carpio* L) on blood indices. *Biochemical and Cellular Archives*. **2020**, 20(2): 5173-5176..
19. Omar Khaled Attallah, Thafer Thabit Mohammed and Nasr Nuri Al-Anbari. Effect of Adding Grape Pomace and Resveratrol on Some Physiological Traits and Gene Expression to Prevent Hemorrhagic



- Fatty Liver Syndrome in Laying Hens . IOP Conference Series: Earth and Environmental Science.2022, 1060 (1), 012076. doi:10.1088/1755-1315/1060/1/0120.
20. Iranpour D, Hassanpour M, Ansari H, Tajbakhsh S, Khamisipour G, Najafi A. Phylogenetic groups of *Escherichia coli* strains from patients with urinary tract infection in Iran based on the new Clermont phylotyping method. *BioMed Res Internat* **2015**; 1-8.
  21. Collee J, Fraser A, Marmion B, Simons A. Mackie and McCartney's Practical Medical Microbiology. 14<sup>th</sup> ed. Churchill Livingstone USA. **1996**; pp: 561.
  22. Brooks GF, Karen C, Carroll MD. Medical Microbiology. Jawetz, Melnick and adelberg's. 24<sup>th</sup> ed. McGraw-Hill Companies, Inc. USA. **2007**; pp: 266-267.
  23. Essa RH, Al-Zubaidi HKA, Rasool KH, Hussein NH. Study of some bacteriological and immunological parameters in urinary tract infection of pregnant women. *Int Educ Res J* **2016**; 2: 2454-9916.
  24. Van Laar TA, Chen T, Childers BM, Chen P, Abercrombie JJ, Leung KP. Genome sequence of a multi-drug-resistant strain of *Klebsiella pneumoniae*, BAMC 07-18, isolated from a combat injury wound. *Genome Announc*, **2014**; 2(6):1230-1244.
  25. Abdalla AE, Shadan AA. Screening for *in vitro* biofilm formation ability of locally isolated uropathogenic *Escherichia coli* (UPEC). *Iraqi J Sci* **2015**; 56(2):1310-1314.
  26. Al-Rubyaie NS. Detection of antiseptic resistant genes in multidrug resistant *Pseudomonas aeruginosa* and *Klebsiella pneumonia* from hospital environment. M.Sc. Thesis. Mustansiriyah University, Iraq. **2021**; pp: 78.
  27. Al-Husseini B. Toxin-Antitoxin system as Anti-persister regulator in *Pseudomonas aeruginosa*. Ph.D. Dissertation. Mustansiriyah University. **2020**; pp: 79-81.
  28. Al-Musawi AM. Genotypic and phenotypic typing of clinical *Klebsiella pneumonia* local isolates. M.Sc. Thesis. Mustansiriyah University, Iraq. **2019**; pp: 22.
  29. Mirzaie A. Ranjbar R. Antibiotic resistance, virulence-associated genes analysis and molecular typing of *Klebsiella pneumonia* strains recovered from clinical samples. *AMB Express* **2021**; 11(1): 1-11.
  30. Ferreira R, Silva B, Rezende1 G, Silva R, Silva A, Campanini1 E, Brito M, Silva E, Freire C, Cunha A, Pranchevicius M. High Prevalence of Multidrug-Resistant *Klebsiella pneumoniae* Harboring Several Virulence and b-Lactamase Encoding Genes in a Brazilian Intensive Care Unit. *Frontiers Microbiol* **2019**; 9, 3198-3205.
  31. Al-Aajem B, Saleem A, Jasim H. Detection of extended spectrum beta-lactamase genes (BLACTX and BLA-shv) and same virulent genes in *K. pneumoniae* isolated from urinary tract infection. *Biochem Cell Arch* **2021**; 21 (2): 4705-4710.
  32. Alsanie W. Molecular diversity and profile analysis of virulence-associated. *Pract Lab Med*. **2020**; 19, 1-12
  33. Remya P, Shanthi M, Sekar U. Characterization of Virulence Genes Associated with Pathogenicity in *Klebsiella pneumonia*. *Indian J Med Microbiol* **2019**; 37(2): 210-218.
  34. Abdulateef SM, Al-Bayar MA, Majid AA, Shawkat SS, Tatar A, Al-Ani MQ. Effect of exposure to different light colors on embryonic development and neurophysiological traits in the chick embryo. *Veterinary World*. 2021 May;14(5):1284.
  35. Z. Al-Fayyadh, D. .; Hasson, A. A. .; Hussein, A. K. .; Hassan, R. K. EFFECT OF HUMIC ACID SPRAY ON GROWTH CHARACTERISTICS OF WHEAT VARIETIES . *Journal of Life Science and Applied Research*. 2020, 1, 10-19.

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