



www.bioinformation.net
Volume 18(5)

Research Article

Received April 2, 2022; Revised May 31, 2022; Accepted May 31, 2022, Published May 31, 2022

DOI: 10.6026/97320630018482

Declaration on Publication Ethics:

The author's state that they adhere with COPE guidelines on publishing ethics as described elsewhere at <https://publicationethics.org/>. The authors also undertake that they are not associated with any other third party (governmental or non-governmental agencies) linking with any form of unethical issues connecting to this publication. The authors also declare that they are not withholding any information that is misleading to the publisher in regard to this article.

Declaration on official E-mail:

The corresponding author declares that lifetime official e-mail from their institution is not available for all authors

License statement:

This is an Open Access article which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly credited. This is distributed under the terms of the Creative Commons Attribution License

Comments from readers:

Articles published in BIOINFORMATION are open for relevant post publication comments and criticisms, which will be published immediately linking to the original article without open access charges. Comments should be concise, coherent and critical in less than 1000 words.

Edited by P Kanguane

Citation: Rafeeq *et al.* Bioinformation 18(5): 482-487 (2022)

Effect of rifampicin combination-regimens against multi-drug resistant strains in North India

Misbahuddin M Rafeeq¹, Alaa Hamed Habib², Ahmad Alzamami³, Norah A. Alturki⁴, Mutaib M. Mashraqi⁵, Youssef Saeed Alghamdi⁶, Saleh Alshamrani⁵, Afaf Awwadh Alharthi⁷ & Suhail Ahmad^{8*}

¹Department of Pharmacology, Faculty of Medicine, Rabigh, King Abdulaziz University, Jeddah, 21589, Saudi Arabia; ²Department of Physiology, Faculty of Medicine, King Abdulaziz University, Jeddah, 21589, Saudi Arabia; ³Clinical Laboratory Science Department, College of Applied Medical Science, Shaqra University, AlQuwayiyah 11961, Saudi Arabia; ⁴Clinical Laboratory Science Department, College of Applied Medical Science, King Saud University, Riyadh 11433, Saudi Arabia; ⁵Department of Clinical Laboratory Sciences, College of Applied Medical Sciences, Najran University, Najran 61441, Saudi Arabia; ⁶Department of Biology, Turabah University College, Taif University, P.O.BOX 11099, Taif 21944, Saudi Arabia; ⁷College of Applied Medical Sciences, Department of Clinical Laboratory Sciences, Taif University, Taif, Saudi Arabia; ⁸Department of Bioengineering, Integral University, Lucknow, India; *Corresponding author:

Author contacts:

Misbahuddin M Rafeeq - Email: marafeeq@kau.edu.sa

Alaa Hamed Habib - Email: ahabib@kau.edu.sa

Ahmad Alzamami - Email: aalzamami@su.edu.sa

Norah A Alturki - Email: noalturki@ksu.edu.sa

Mutaib M Mashraqi - Email: mmmashraqi@nu.edu.sa

Youssef Saeed Alghamdi - Email: ysghamdi@tu.edu.sa

Saleh Alshamrani - Email: saalshamrani@nu.edu.sa

Afaf Awwadh Alharthi - Email: a.awwadh@tu.edu.sa

Suhail Ahmad - Email: ssbiotechnologist@gmail.com & suhailamd@iul.ac.in

Abstract

It is well-acknowledged that 'combination therapy' of antibiotics is indispensable for the treatment of patients suffering from serious bacterial infections. Therefore, it is of interest to collect data from the *in vitro* tests using 'rifampicin-cefotaxime' and 'rifampicin-tetracycline' combination regimens against multi drug resistant *Escherichia coli* and *Klebsiella pneumoniae* strains of nosocomial source in order to determine the effectiveness of the combination therapy. The minimum inhibitory concentration (MIC) values for cefotaxime, tetracycline and rifampicin antibiotics were found to be comparatively high for each of the antibiotics when given individually. However, carefully prepared combination-regimens exhibited significant inhibitory effect on the same bacterial isolates. DNA fragmentation study confirmed that 'rifampicin-cefotaxime' and 'rifampicin-tetracycline' combination-regimens could cause breakage of the bacterial DNA. Thus, we show that combination-regimens namely, 'rifampicin-cefotaxime' and 'rifampicin-tetracycline' were found to be capable of maintaining rifampicin susceptibility in the *E. coli* and *K. pneumoniae* strains. However, this susceptibility was not maintained by only rifampicin. More data using animal model experiments are needed for confirming and deriving translational benefits from these findings in future.

Key words: Antibiotic combinations; Cefotaxime; DNA fragmentation assay; *Escherichia coli*.

Background:

Infectious diseases occurring due to multidrug resistant Gram-negative bacteria are a major therapeutic challenge [1, 2] in community as well as hospital settings [3]. If we consider the population living in the developed nations, the average life tenure for an individual has increased. Albeit this augmented tenure is also coupled to many other health issues because of steep upsurge in the cases of obesity and diabetes. Accordingly, this has resulted in the rise of 'chronic wound-infections'. In USA, such infections had been estimated to touch some 6.5×10^6 individuals and it was projected to produce a burden of 25×10^9 US\$ per annum, along with a high chance of further increase in the years to come [4]. In order to overcome/decrease the pace of drug-resistance, disease treatment employing 'combination-regimens' of carefully selected antibiotics is a routine practice in many health care centers [5-7]. This type of approach could lead to a sort of synergism thereby resulting in overall enhanced efficiency of the treatment as well as decrease in the total amount of each drug consumed. Moreover, the chance of many contra-indications could also be minimized. Most importantly, this could duly decrease the total expenditure incurred [8, 9]. Furthermore, it had been argued that using combination-regimens i.e. including more than one antibiotic, each having a different 'action-mode' and/or 'target' might reduce the chance of emerging drug-resistance while the treatment continued [10]. As far as the case of chronic infections is considered, it is of particular importance, as the drug-therapy associated with such treatment may be significantly prolonged. A couple of substitute approaches to increase the effectiveness of known antibiotic treatments against hardier bacteria had been also suggested. For instance, augmenting the drug-toxicity in a synergistic style by combining these antibiotics to other chemicals or to metals possessing dissimilar targets is quite a germane approach. This line of attack on bacteria expressively decreases the chance of appearance of bacteria which could concurrently resist both of the anti-bacterial [11]. Therefore, it is of interest to document data from the *in vitro* tests using 'rifampicin-cefotaxime' and 'rifampicin-

tetracycline' combination regimens against multi drug resistant *Escherichia coli* and *Klebsiella pneumoniae* strains of nosocomial source in order to determine the effectiveness of the combination therapy.

Materials and Methods:

Bacterial isolates:

The bacterial strains (*E. coli* and *K. pneumoniae*) were obtained from cultured urines on selective media. We used the reference methods described by Cowan and Steel [12] to isolate and identify the bacterial strains.

Broth-dilution test:

We used an adapted protocol that was based on the famous macro broth dilution protocol as described by Ibrahim *et al.* [13] to establish the values for minimum inhibitory-concentration (MIC). Bacterial culture which was incubated for 12 hours in nutrient broth was diluted 100 folds using sterile nutrient broth (100 μ l of bacterial-cultures added to 10ml of nutrient broth that had 10^8 cfu of bacteria). Progressively, we added increasing volumes of different concentration of antibiotics and their combinations to the test-tubes having the bacterial-cultures to measure the inhibiting-concentration in a particular tube inhibiting the bacterial-growth. These test-tubes were incubated at 37°C for 18-24 hours. We inspected these test-tubes for visible turbidity while the O.D. values were measured at a wavelength of 620nm. Sterile nutrient broth was used as a control. The minimum concentration which inhibited the visible-growth of the bacterial strains was noted as the corresponding MIC-value. Further, MICs for the selected antimicrobials were tested by HiComb MIC test strips (Hi-media, India).

Agar well-diffusion test:

Agar well-diffusion test was used to assess the antibacterial effects of the antibiotics and their combinations [14]. Nutrient-agar was duly added to all of the Petri-plates which had a diameter of 9 cm.

After the solidification of the agar, pertinent number of wells, each having a 5 mm diameter, were prepared in the agar plate(s). 100µl (10⁵cfu) of each of the diluted bacterial culture was inoculated onto the nutrient agar-plates with the aid of sterile cotton-swabs. Rifampicin, tetracycline, cefotaxime as well as their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) were added to each of the bored wells in the agar plates. These were allowed to diffuse at normal room temperature for about 20 minutes. Post-incubation at 37°C for 24h, all of the plates were scrutinized for 'growth inhibition-zones' along with duly measuring the respective zone-diameters.

DNA fragmentation test:

DNA fragmentation assay was carried out under stress conditions at different concentrations of cefotaxime, tetracycline and rifampicin and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) employing the method of Fernandez *et al.* [15].

Table 1: Minimum inhibitory concentration of rifampicin, cefotaxime, tetracycline and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) against bacterial strains

Bacterial strain	Minimum Inhibitory Concentration (mg/L)				
	CTX	TET	RIF	RIF+CTX	RIF+TET
<i>E. coli</i> (E2)	15.62	7.81	31.25	1.95	0.97
<i>E. coli</i> (E3)	15.62	7.81	31.25	1.95	0.97
<i>E. coli</i> (E1)	15.62	31.25	7.81	0.97	1.95
<i>K. pneumoniae</i> (K1)	31.25	7.81	15.25	0.97	1.95
<i>K. pneumoniae</i> (K2)	15.62	31.25	7.81	1.95	3.9
<i>K. pneumoniae</i> (K3)	7.81	15.62	31.25	1.95	3.9

*The data refer to mean value of three replicates. RIF= Rifampicin, CTX = Cefotaxime, TET= Tetracycline

Table 2: MICs of antibiotics used against *E. coli* and *K. pneumoniae*

Antibiotics	MIC (mg/L)					
	<i>E. coli</i> isolates			<i>K. pneumoniae</i> isolates		
	E1	E2	E3	K1	K2	K3
Aztreonam	>2	>2	>2	>2	1	>2
Amikacin	8	4	8	8	8	4
Amoxycillin	>240	240	>240	>240	>240	240
Ceftazidime	256	256	64	256	>256	256
Cefepime	128	128	32	32	128	64
Ceftriaxone	128	128	>256	256	32	256
Ciprofloxine	2	>2	1	>2	2	2
Streptomycin	10	7.5	7.5	5	5	10
Erythromycin	60	60	30	60	30	30
Amoxyclav	30	60	30	30	10	60

Results and Discussion:

Minimum inhibitory concentration (MIC) determination:

Antibacterial activity of cefotaxime, tetracycline and rifampicin and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) were evaluated against Gram-negative (*E. coli* and *K. pneumoniae*) bacterial strains. MIC measurement was performed employing an adapted version of the famous macro-broth dilution protocol [13]. MIC values of cefotaxime, tetracycline and rifampicin were found to be individually high for *E. coli* and *K. pneumoniae* strains. The MIC-values of cefotaxime, tetracycline and rifampicin for *E. coli* (E1) were found to 15.625, 31.25, and 7.81 mg/L, respectively, whereas the same for *K. pneumoniae* (K1) were determined to be 31.25, 7.81, 15.62 mg/L, respectively. In contrast

to this, the MIC value of rifampicin combined with cefotaxime or tetracycline was found to be comparatively low for *E. coli* and *K. pneumoniae* strains (Table 1). MIC-values corresponding to 'rifampicin-cefotaxime' and 'rifampicin-tetracycline' combinations for *E. coli* (E1) were found to be 0.97 and 1.95 mg/L, respectively, whereas the same for *K. pneumoniae* (K1) were found to be 0.97 and 1.95 mg/L, respectively. Our results are consistent with the findings in another study where the author has described results of combination therapy of rifampicin with other antibiotics against multi drug resistant *Pseudomonas aeruginosa* isolates [16]. Furthermore, a combination of rifampicin with penicillin or ampicillin was established to exhibit a full synergistic bactericidal activity way back in 1982 by Tuazon *et al.* [17]. Further, the MICs of the different antibiotics were determined on both *E. coli* and *K. pneumoniae* isolates as presented in table 2 (Figure 1). Very high MICs were obtained for amoxycillin, ceftazidime, cefepime, erythromycin and ceftriaxone indicating that the studied strains were highly resistant to these antibiotics. The MICs of aztreonam, amikacin, ciprofloxine and streptomycin were moderate, but still in the resistant range. The MIC values obtained against different antibiotics in our study were in due agreement with earlier reports [18-20].

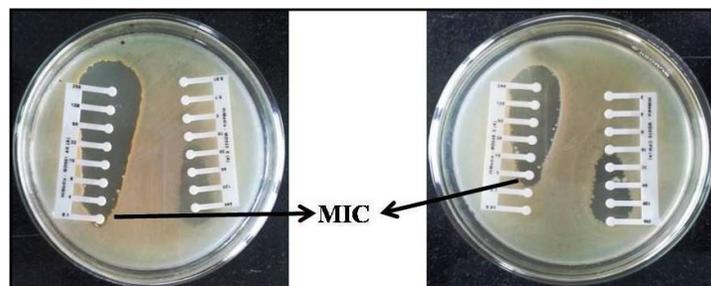


Figure 1: A typical HiComb MIC test (Himedia, India). The base-point of the balloon shaped region (marked with an arrow) corresponds to MIC of the test antibiotic.

Determination of antimicrobial effect:

The 'Zone of Inhibition'-tests revealed that all of the bacterial isolates (nosocomial strains of *E. coli* and *K. pneumoniae*) were not susceptible to the solo drugs i.e. rifampicin, cefotaxime and tetracycline, when given individually (Table 3). However, the antibiotic combination regimens exhibited significant inhibitory effect on *E. coli* as well as *K. pneumoniae* isolates. The zones of inhibition corresponding to 'rifampicin-cefotaxime' and 'rifampicin-tetracycline' combinations against *E. coli* (E1) were found to possess diameters of 20 and 17 mm, respectively; while the same for *K. pneumoniae* (K2) were found to have diameters of 19 and 17 mm, respectively. Therefore, it can be concluded that 'rifampicin-cefotaxime' and 'rifampicin-tetracycline' combinations displayed potent antimicrobial activity against the tested isolates. The zones of inhibition shown by cefotaxime, tetracycline, rifampicin and their combinations (rifampicin-cefotaxime and cefotaxime-tetracycline) on *E. coli* and *K. pneumoniae* strains are shown in Figure 2 and Figure 3. In a notable study which supports the findings described herein, the authors have reported that the

combination of rifampicin with other selected antibiotics demonstrated full synergistic bactericidal activity [17].

Table 3: Zones of inhibition by cefotaxime, tetracycline and rifampicin and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) on bacterial strains.

S. No.	Bacterial strain	Zone of inhibition (mm)				
		RIF	CTX	TET	RIF+CTX	RIF+TET
1.	<i>E. coli</i> (E1)	R	RR	20	17	
2.	<i>E. coli</i> (E2)	R	RR	19	18	
3.	<i>E. coli</i> (E3)	R	RR	16	17	
4.	<i>K. pneumoniae</i> (K1)	R	RR	17	16	
5.	<i>K. pneumoniae</i> (K2)	R	RR	19	17	
6.	<i>K. pneumoniae</i> (K3)	R	RR	17	18	

*The data is mean of three replicates; CTX = Cefotaxime, TET = Tetracycline, RIF = Rifampicin, R = Resistant

Rifampicin has a number of characteristics that might make it significantly effective when used in combination with other antibiotics, namely, its potent bactericidal activity [21], modest activity against non-growing cells [22], ability to penetrate the cells [23] and a variety of tissues and compartments, such as cerebrospinal fluid and bone [24]. Hence, our data supports the idea of designing 'rifampicin-cefotaxime' as well as 'rifampicin-tetracycline' combination regimens for the effective treatment of multidrug-resistant strains of *E. coli* and *K. pneumoniae* of clinical origin.



Figure 2: Zone of inhibition shown by rifampicin, cefotaxime, tetracycline and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) on *E. coli*.

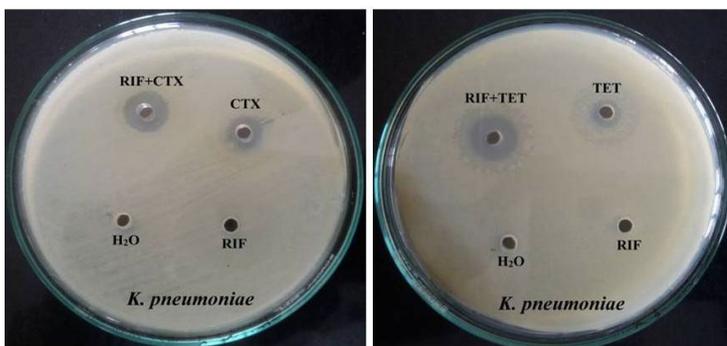


Figure 3: Zone of inhibition shown by rifampicin, cefotaxime, tetracycline, and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) on *K. pneumoniae*.

DNA fragmentation assay:

DNA fragmentation study was carried out using rifampicin, cefotaxime, tetracycline and their combinations ('rifampicin-cefotaxime' and 'rifampicin-tetracycline') on DNA of the aforementioned bacterial isolates. We observed that rifampicin-cefotaxime and rifampicin-tetracycline combinations are lethal for bacterial DNA as shown in **Figure 4** and **Figure 5**. In fact, rifampicin acts by inhibiting the DNA-dependent-RNA-polymerase, thereby preventing expression of the bacterial genes [25]. However, in the current study some extent of DNA fragmentation was also observed in cefotaxime and tetracycline treated isolates. Accordingly, Molina-Quiroz *et al.* [26] studied synergistic effect of tellurite/cefotaxime in *E. coli*. They showed that the tellurite/cefotaxime treatment caused cellular damage. Increased levels of intracellular superoxide and OH[•] produced by tellurite and cefotaxime, respectively, generated direct damage to DNA [26]. In contrast to this, tetracycline has been shown to induce SOS response in *Vibrio cholera* [27]. The SOS response is an inducible pathway governing DNA repair that was first described in *E. coli* [28].

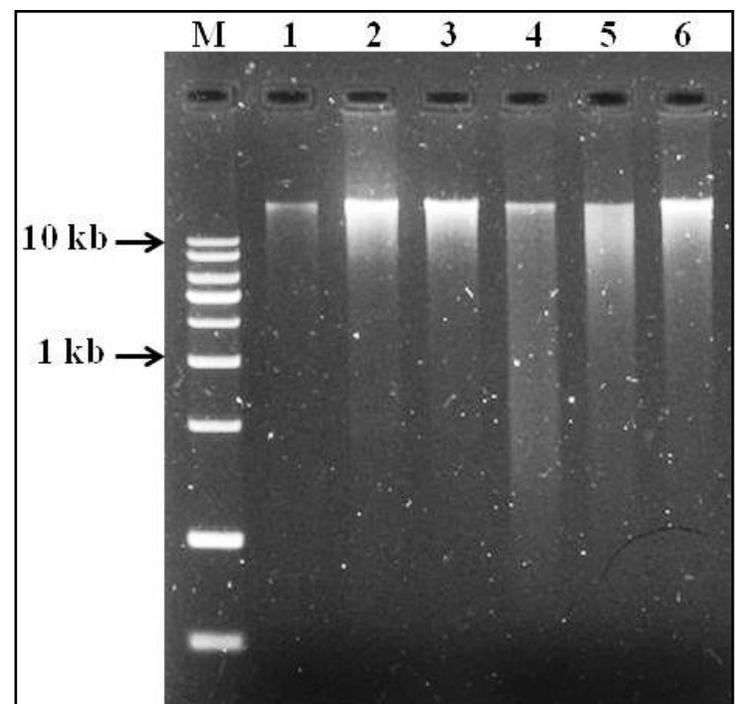


Figure 4: DNA fragmentation showed by cefotaxime, tetracycline, rifampicin and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) on *E. coli*. M - Marker; Lane1 - Control; Lane2 - Cefotaxime; Lane3 - Tetracycline; Lane4 - Rifampicin; Lane5 - Rifampicin + Cefotaxime; Lane6 - Rifampicin + Tetracycline

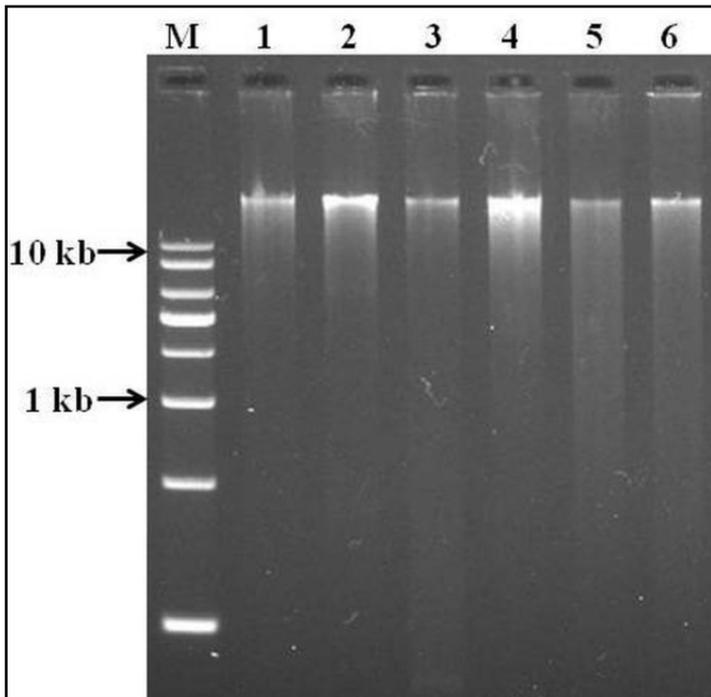


Figure 5: DNA fragmentation showed by rifampicin, cefotaxime, tetracycline and their combinations (rifampicin-cefotaxime and rifampicin-tetracycline) on *K. pneumoniae*. M - Marker; Lane1 - Control; Lane2 - Cefotaxime; Lane3 - Tetracycline; Lane4 - Rifampicin; Lane5 - Rifampicin + Cefotaxime; Lane 6 - Rifampicin + Tetracycline

Fragmentation of the Chromosomal-DNA might directly or indirectly relate to the phenomenon of death of the cell. As opposed to DNA-fragmentation concerning higher eukaryotic-cells, fragmentation of the same in microbes has been explored to somewhat lesser extent. In fact, Chromosomal fragmentation of DNA of chromosomal origin might represent cell-death because this in turn is the outcome of substantial breaks in the double strand of the DNA. In higher-cells, the said observation might be a sequel to an active 'programmed-cell-death/apoptosis', where the DNA is cleaved via an activated endonuclease [29]. Else, this fragmentation could also occur passively by necrotic cell-death. However, the passive-type fragmentation of the DNA occurs more frequently in the microbes that get destroyed due to varied reasons. Nevertheless, researches indicate a feasibility of occurrence of 'programmed cell death/apoptosis' in single-celled bacterial pathogens as well [30]. In actual fact, bactericidal-drugs might activate an apoptosis like pathway. Scientists have noted that bactericidal-drugs could trigger the formation of hydroxyl-radicals which in turn could underpin cell death. It has been argued that tolerant-type bacteria, that are somewhat resistant to certain bactericidal-drugs, might actually be the bacteria that possess a disabled apoptotic pathway/program [31].

Conclusion:

Combination-regimens of 'rifampicin-cefotaxime' and 'rifampicin-tetracycline' were found to be capable of maintaining rifampicin

susceptibility in the *E. coli* and *K. pneumoniae* strains unlike solo rifampicin. More data from animal model experiments are needed for confirming and deriving translational benefits from these findings in future.

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

References:

- [1] Shaikh S *et al.* *Current Drug Metabolism* 2015 **16**:362. [PMID: 26419545]
- [2] Shaikh S *et al.* *Saudi journal of biological sciences* 2015 **22**:90. [PMID: 25561890]
- [3] Shaikh S *et al.* *Lett Appl Microbiol* 2016 **62**:419. [PMID: 26997253]
- [4] Sen CK *et al.* *Wound Repair Regen* 2009 **17**:763. [PMID: 19903300]
- [5] Susanto BO *et al.* *Clin Pharmacol Ther* 2020 **108**:274 [PMID: 32080839]
- [6] Rao GG *et al.* *Clin Microbiol Infect* 2018 **24**:689 [PMID:29269090]
- [7] Tang HJ *et al.* *Antimicrob Agents Chemother* 2013 **57**:5717 [PMID: 23959320]
- [8] Boyd N & Nailor MD. *Pharmacotherapy* 2011 **31**:1073. [PMID: 22026395]
- [9] El-Hafi B *et al.* *J Infect Dev Ctries* 2018 **12**:14. [PMID: 31804989]
- [10] Rahal JJ. *Clin Infect Dis* 2006 **43**:95. [PMID: 16894522]
- [11] Mitchell G *et al.* *J Antimicrob Chemother* 2012 **67**:559. [PMID: 22129590]
- [12] Cowan SF & Steel KJ. *Manual for the Identification of the Medical Bacteria*, Cambridge, UK Cambridge University Press, 1970.
- [13] Ibrahim MB *et al.* *J Pharm Res Dev* 1997 **2**:20.
- [14] Okeke MI *et al.* *J Ethnopharmacol* 2001 **78**:119. [PMID: 11694355]
- [15] Fernández JL *et al.* *Appl Environ Microbiol* 2008 **74**:5925. [PMID: 18689511]
- [16] Hu YF *et al.* *BMC Infect Dis* 2016 **16**:444. [PMID: 27553962]
- [17] Tuazon CU *et al.* *Antimicrob Agents Chemother* 1982 **21**:525. [PMID: 6808911]
- [18] Shaikh S *et al.* *Iranian Journal of Science & Technology* 2017 **41**:1011 [https://doi.org/10.1007/s40995-017-0340-8].
- [19] Faheem M *et al.* *PLoS One* 2013 **8**:e56926. [PMID: 23437273]
- [20] Shaikh S *et al.* *Saudi Journal of Biological Sciences* 2015 **22**:37. [PMID: 25561881]
- [21] Maltempe FG *et al.* *Tuberculosis (Edinb)* 2017 **104**:24. [PMID: 28454646]
- [22] Sala C *et al.* *Antimicrob Agents Chemother* 2010 **54**:4150. [PMID: 20679505]
- [23] Dartois V. *Nature reviews Microbiology* 2014 **12**:159. [PMID: 24487820]
- [24] Spellberg B & Lipsky BA. *Clin Infect Dis* 2012 **54**:393. [PMID: 22157324]
- [25] Hartmann G *et al.* *Biochim Biophys Acta* 1967 **145**:843. [PMID: 4863911]

- [26] Molina-Quiroz RC *et al.* *PLoS ONE* 2013 **8**:e79499. [PMID: 24260236]
- [27] Baharoglu Z and Mazel D, *Antimicrob Agents Chemother* 2011 **55**:2438. [PMID: 21300836]
- [28] Butala M *et al.* *Cell Mol Life Sci* 2009 **66**:82. [PMID: 18726173]
- [29] Nagata S. *Exp Cell Res* 2000 **256**:12. [PMID: 10739646]
- [30] Tanouchi Y *et al.* *Trends in microbiology* 2013 **21**:265. [PMID: 23684151]
- [31] Valenti A *et al.* *Nucleic Acids Res* 2006 **34**:2098. [PMID: 16617150]

