

## Determination of Tigecycline and Colistin Susceptibility among Carbapenemase Producing Isolates of Gram Negative Bacteria in a Tertiary Care Hospital

Justine Auxilia Irene Chinnappan\*, Lakshmi Priya N, Umadevi U, Thasneem Banu S

Institute of Microbiology, Madras Medical College, Chennai, India

\*Corresponding author: Justine Auxilia Irene Chinnappan, Institute of Microbiology, Madras Medical College, Chennai, India

**Citation:** Chinnappan JAI, Lakshmi Priya N, Umadevi U, Thasneem BS (2020) Determination of Tigecycline and Colistin Susceptibility among Carbapenemase Producing Isolates of Gram Negative Bacteria in a Tertiary Care Hospital. J Microbiol Genet 05: 128. DOI: 10.29011/2574-7371.100028

**Received Date:** 15 October, 2020; **Accepted Date:** 12 November, 2020; **Published Date:** 18 November, 2020

### Abstract

**Introduction:** Management of infections with carbapenemase producing Gram negative bacteria has become challenging. Colistin and tigecycline are the drug considered for the treatment of these multidrug resistance bacterial infections. This study was aimed to determine the colistin and tigecycline susceptibility among carbapenemase producing Gram negative bacteria.

**Materials and Methods:** This study was conducted in Institute of Microbiology, Madras Medical College, Chennai with 75 carbapenemase producing Gram negative bacteria from various clinically relevant samples. Bacterial identification was done by standard microbiological protocol and the carbapenemase production was confirmed by Modified Hodge Test. The Colistin and Tigecycline MIC value were determined by E-strip (HIMEDIA) method and interpreted using EUCAST-BSAC 2017 guidelines.

**Result:** Among the 75 isolates included in the study, 26.6% were *Acinetobacter* spp, 22.7% *Klebsiella pneumoniae*, 18.7% *E.coli*, 16% *Klebsiella oxytoca* and 16% *Pseudomonas* spp. The Colistin MIC<sub>50</sub> was 1 µg/ml and 0.75 µg/ml for *Enterobacteriaceae* and Non-Fermenting Gram Negative Bacilli (NFGNB) respectively whereas colistin MIC<sub>90</sub> was 1.5 µg/ml for both *Enterobacteriaceae* and NFGNB. Both the colistin MIC<sub>50</sub> and MIC<sub>90</sub> for *Enterobacteriaceae* and NFGNB were found to be susceptible. The Tigecycline MIC<sub>50</sub> and MIC<sub>90</sub> were 0.094 µg/ml and 0.5 µg/ml respectively for both *Enterobacteriaceae* and NFGNB. Tigecycline MIC<sub>90</sub> for NFGNB was found to be resistant. One among the 75 carbapenemase producing gram negative bacilli tested i.e *Acinetobacter* spp was resistant to colistin with MIC of 8µg/ml. Two *Klebsiella oxytoca* showed intermediate susceptibility and six *Acinetobacter* spp showed resistance to tigecycline. The *Acinetobacter* spp. which was resistant to colistin was also resistant to tigecycline with MIC of 0.75 µg/ml.

**Conclusion:** The study showed the frequency of colistin and tigecycline resistance to be 1.3% and 12.5% respectively. With these findings we conclude that colistin and tigecycline may be considered for the treatment of Carbapenemase producing Gram negative bacterial infections. However surveillance studies are recommended to organise treatment protocol and determine the dosage based on these surveillance data.

**Keywords:** Carbapenemase producing Gram negative bacteria; Colistin; MIC; Tigecycline

### Introduction

In this era of worsening antimicrobial resistance, gram negative bacteria resistant to carbapenem poses a special clinical challenge because they have been the most active and potent agents for a longtime against multidrug resistant gram negative bacteria.

This is further supported by World Health Organization priority pathogens list for research and development of new antibiotics in 2017 which enlists carbapenem-resistant *Enterobacteriaceae* (CRE), carbapenem-resistant *Acinetobacter baumannii*, and carbapenem-resistant *Pseudomonas aeruginosa* as critical priority [1].

The mechanisms responsible for carbapenem resistance including porin mutations, efflux pump upregulation and

carbapenemase production are explained in detail by Nordmann and Poirel [2]. Of significance are chromosomally encoded porin gene mutation (such as OprD), overexpression of genes encoding for efflux pumps (such as MexAB-OprM, MexXY-OprM, or MexCD-OprJ) and the carbapenemases like KPC, VIM, IMP, NDM and OXA-48 types [2-4]. Resistance due to carbapenemase production is clinically most important as they not only hydrolyze almost all beta-lactams and confer high Minimum Inhibitory Concentration (MICs) but are also encoded by genes transferable horizontally by plasmids or transposons which also have genes encoding for other resistance determinants [5].

Reports of carbapenemase-mediated resistance continue to increase globally [6,7], leading clinicians to resort to other antibiotics with therapeutic success against carbapenemase-producing Gram-negative bacteria. Currently, colistin (polymyxin E) and tigecycline have become the antibiotics of last resort for carbapenemase-producing Gram-negative bacteria [8,9]. Increasing colistin and tigecycline consumption has been concurrent with increasing reports of tigecycline and colistin resistance, especially during therapy. Already, clinical outbreaks involving colistin resistant KPC-producing *Klebsiella pneumoniae* have been reported in the United States and Italy with worrying recurrence leaving very limited treatment options [5,6]. Increasing nonsusceptibility to colistin and tigecycline during colistin and tigecycline monotherapy has engendered the use of colistin and/or tigecycline in double and triple combinations with a carbapenem, an aminoglycoside (amikacin, gentamycin and tobramycin), rifampicin, fosfomycin or fluoroquinolone [8,9].

In view of the above, this study was conducted to isolate and identify the carbapenemase producing Gram negative bacteria from patients admitted in various wards and determine their colistin and tigecycline susceptibility pattern.

## Materials and Methods

### Study Design

This cross-sectional study was carried out in the Institute of Microbiology Madras Medical College and Rajiv Gandhi Government General Hospital, Chennai. The study was conducted for a period of 6 months from (April 2017 to September 2017)

**Sample Size:** 75 carbapenemase producing Gram negative bacteria

### Inclusion Criteria

- Non-repetitive isolates of Carbapenemase producing Gram negative bacterial isolates.
- Only one isolate per sample in polymicrobial infection were included.

## Bacterial Identification and Phenotypic Analysis of Carbapenems

Non-repetitive clinically relevant isolates of Gram negative bacteria from various clinical samples including blood, sputum, pus, wound swabs, bronchial wash, tracheal aspirates and urine were collected. The isolates were identified as per Standard Microbiological protocol. Antimicrobial susceptibility testing of the isolates were done by Kirby Bauer's disc diffusion technique on Mueller-Hinton agar, as per Clinical Laboratory Standard Institute (CLSI) guidelines 2017. Isolates with intermediate susceptibility or resistance to meropenem and imipenem by disc diffusion method were screened for the production of carbapenemase.

### Phenotypic Detection of Carbapenemases

A phenotypic detection of carbapenemases was done by Modified Hodge test.

**Modified Hodge Test (MHT) [10]:** 0.5 McFarland standard suspension of *E.coli* ATCC 25922 was prepared in saline and diluted 1:10 in saline. A lawn culture of 1:10 dilution *E.coli* ATCC 25922 was done on to a Mueller Hinton Agar plate and allowed to dry for 3-5 minutes. A 10µg meropenem disc is placed in the center of the test area. In a straight line, the test organism was streaked from the edge of the disc to the edge of the plate and incubated at 37°C for 16-20 hours. Positive and negative controls were included.

### Interpretation

Enhanced growth (Clover-leaf indentation) = Positive

No enhanced growth = Negative

### Colistin Susceptibility Test

Colistin MIC was determined by Epsilon Meter Test (E-test) using colistin E-strips of concentration gradient 0.016 to 256µg/ml (HIMEDIA) as per the EUCAST guidelines 2017 [11]. *E.coli* ATCC 25922 was used for Quality control (QC range, 0.25-2µg/ml).

Four or five identical colonies were inoculated into 5ml of saline and incubated for 2 hours to match 0.5 McFarlands standards. Lawn culture of the suspension was made onto Mueller Hinton Agar plate. The E strip was placed on the Mueller Hinton Agar plate using a sterile applicator and incubated at 37°C for 20-24 hrs. After incubation whereby bacterial growth becomes visible, an elliptical zone of inhibition correlating with the gradient concentration of antibiotic. The MIC value is read from the scale in terms of µg/ml where the point of ellipse intersects the strip (Table 1).

Organism	Minimum Inhibitory Concentration (MIC) - µg/ml		
	Sensitive	Intermediate	Resistant
<i>Enterobacteriaceae</i>	≤2	-	>2
<i>Pseudomonas spp.</i>	≤2	-	>2
<i>Acinetobacter spp.</i>	≤2	-	>2

**Table 1:** MIC interpretive criteria for Colistin [11].

### Tigecycline Susceptibility Test

*Pseudomonas* species is intrinsically resistant to Tigecycline, hence *Enterobacteriaceae* and *Acinetobacter spp.* were tested for Tigecycline MIC by using tigecycline E-strips of concentration gradient 0.016 to 256µg/ml (HIMEDIA) as per the EUCAST guidelines 2017. *E.coli* ATCC 25922 was used for Quality control (QC range, 0.03-0.25µg/ml) (Table 2).

Organism	Minimum Inhibitory Concentration (MIC) - µg/ml		
	Sensitive	Intermediate	Resistant
<i>Enterobacteriaceae</i>	≤1	>1 to ≤2	>2
<i>Acinetobacter spp.</i>	≤0.25	-	≥0.5

**Table 2:** MIC interpretive criteria for Tigecycline [11,12].

**Statistical Analysis:** Data were entered in Microsoft Excel 2013. MIC<sub>50</sub> and MIC<sub>90</sub> were calculated for colistin and Tigecycline among various organisms. MIC<sub>50</sub> and MIC<sub>90</sub> were defined as lowest concentration of antibiotics (lowest MIC) at which 50% and 90% of the tested strains were inhibited.

### Results

A total of 112 Carbapenem resistant gram negative isolates by disc diffusion were screened for Carbapenemase production by Modified Hodge Test, out of which 75 isolates tested positive. Clinical samples for the isolates included in the study and distribution of isolated bacteria are provided in (Table 3).

Clinical samples	<i>E. coli</i>	<i>K.pneumoniae</i>	<i>K.oxytoca</i>	<i>Pseudomonas aeruginosa</i> & other <i>Pseudomonas spp.</i>	<i>Acinetobacter spp.</i>	Total
Urine	11	8	7	5	7	38
Pus	2	4	3	5	6	20
Tracheal aspirate	1	3	2	2	1	9
Sputum	0	2	0	0	2	4
Body fluids	0	0	0	0	2	2
Bronchial wash	0	0	0	0	1	1
Blood	0	0	0	0	1	1
Total	14	17	12	12*	20	75

\*Out of 12 *Pseudomonas spp.* 7 were *Pseudomonas aeruginosa*

**Table 3:** The source and distribution of isolates in clinical samples (n=75).

Among the 75 Modified Hodge Test positive (Carbapenemase producers) isolates, 50.6% isolates were from urine sample and the most frequently isolated organism was *Acinetobacter spp* (26.6%) followed by *Klebsiella pneumoniae*, *E.coli*, *Klebsiella oxytoca* and *Pseudomonas spp*.

Susceptibility pattern of the isolates to colistin and tigecycline is provided in (Table 4).

Organism	Colistin(n=75)		Tigecycline(n=63)		
	S	R	S	I	R
<i>E.coli</i>	14	0	14	0	0
<i>K.pneumoniae</i>	17	0	17	0	0
<i>K.oxytoca</i>	12	0	10	2	0
<i>Pseudomonas aeruginosa</i> & other <i>Pseudomonas</i> species	12	0	-*	-*	-*
<i>Acinetobacter spp.</i>	19	1	14	-	6
Total	74	1	55	2	6

\**Pseudomonas spp.* is intrinsically resistant to Tigecycline hence Tigecycline MIC was not performed for them.

**Table 4:** Determination of the MIC values of the antibiotics tested for the isolates.

Among the 75 carbapenemase producing gram negative bacilli tested, one isolate of *Acinetobacter spp.* (*Acinetobacter baumannii*) from the pus sample of pyothorax was resistant to colistin with MIC of 8µg/ml. Two *Klebsiella oxytoca* showed intermediate susceptibility and six *Acinetobacter spp.* showed resistance to tigecycline. The *Acinetobacter spp.* which was resistant to colistin was also resistant to tigecycline with MIC of 0.75 µg/ml.

The distribution of colistin MIC among the carbapenemase producing gram negative bacteria is shown in (Table 5).

Organism	Number of isolates with colistin MIC (µg/ml)					Total
	0.75	1	1.5	2	8	
<i>E.coli</i>	4	8	2	0	0	14
<i>K.pneumoniae</i>	9	6	1	1	0	17
<i>K.oxytoca</i>	5	6	1	0	0	12
<i>Pseudomonas aeruginosa</i> & other <i>Pseudomonas</i> species	8	2	2	0	0	12
<i>Acinetobacter spp.</i>	10	7	2	0	1	20
Total	36	29	8	1	1	75

**Table 5:** Distribution of colistin minimum inhibitory concentrations in carbapenemase producing Gram negative bacteria (n=75). Maximum isolates (36/75) showed colistin MIC of 0.75 µg/ml, followed by 1 µg/ml(29/75).

The distribution of tigecycline MIC among the carbapenemase producing gram negative bacteria is shown in (Table 6).

Organism	Number of isolates with tigecycline MIC (µg/ml)											Total
	0.047	0.064	0.094	0.125	0.25	0.32	0.5	0.75	1	1.5	2	
<i>E.coli</i>	1	2	3	2	2	1	3	0	0	0	0	14

<i>K.pneumoniae</i>	6	4	2	1	1	0	3	0	0	0	0	17
<i>K.oxytoca</i>		3	2	0	1	0	3		1	1	1	12
<i>Acinetobacter spp.</i>	3	5	3	1	2	0	4	2	0	0	0	20
Total	10	14	10	4	6	1	13	2	1	1	1	63

**Table 6:** Distribution of tigecycline minimum inhibitory concentrations in carbapenemase producing Gram negative bacteria (n=63). Maximum isolates (14/63) showed tigecycline MIC of 0.064 µg/ml, followed by 0.5 µg/ml(13/63).

The colistin and tigecycline MIC<sub>50</sub> and MIC<sub>90</sub> was calculated for *Enterobacteriaceae* and non-fermenters separately are listed in (Table 7).

Antibiotic	Organism	MIC Range	MIC <sub>50</sub> (µg/ml)	Interpretation*	MIC <sub>90</sub> (µg/ml)	Interpretation*
Colistin	<i>Enterobacteriaceae</i>	0.75-2	1	Susceptible	1.5	Susceptible
	NFGNB	0.75-8	0.75	Susceptible	1.5	Susceptible
Tigecycline	<i>Enterobacteriaceae</i>	0.047-2	0.094	Susceptible	0.5	Susceptible
	NFGNB	0.047-0.75	0.094	Susceptible	0.5	Resistant
*The interpretation of the MIC <sub>50</sub> and MIC <sub>90</sub> of all antibiotics tested for all pathogens was performed using the EUCAST-BSAC 2017 guidelines. NFGNB- Non-fermenting Gram negative bacilli						

**Table 7:** Determination and interpretation of the MIC values of the antibiotics tested (n=75).

The Colistin MIC<sub>50</sub> was determined to be 1 µg/ml and 0.75 µg/ml for *Enterobacteriaceae* and NFGNB respectively whereas colistin MIC<sub>90</sub> was determined to be 1.5 µg/ml for both *Enterobacteriaceae* and NFGNB. Both the colistin MIC<sub>50</sub> and MIC<sub>90</sub> for *Enterobacteriaceae* and NFGNB was found to be in the susceptible.

The Tigecycline MIC<sub>50</sub> and MIC<sub>90</sub> were determined to be 0.094 µg/ml and 0.5 µg/ml respectively for both *Enterobacteriaceae* and NFGNB.

## Discussion

The knowledge of antibiotics susceptibility pattern of the bacteria is necessary to overcome the problem of treating infections with resistant isolates. Available treatment options for infection caused by carbapenemase producing GNB are limited currently with colistin and tigecycline being the primary treatment. Increasing use of colistin and tigecycline for carbapenemase producing GNB infections has led to the emergence of colistin and tigecycline resistance in several countries worldwide. The prevalence of this emerging resistance varies between regions based on the availability and use of these antibiotics [13]. Hence this study was aimed to determine the colistin and tigecycline susceptibility pattern in Carbapenemase producing Gram negative bacteria isolated from clinical samples in a tertiary care hospital, Tamil Nadu.

A recent Global surveillance program study involving 39

countries, revealed 88% and 91.1% of carbapenemase producing *Enterobacteriaceae* were susceptible to colistin and tigecycline respectively [14] and a study conducted in Bangalore by Chandran, et al revealed 100% and 94.05% of carbapenemase producing *Enterobacteriaceae* were susceptible to colistin and tigecycline respectively [15]. While in the present study, the colistin and tigecycline susceptibility among *Enterobacteriaceae* were 100% and 94.43% respectively, which indicates lower prevalence of resistance comparatively.

The MIC<sub>50</sub> and MIC<sub>90</sub> for colistin and Tigecycline among *Enterobacteriaceae* in the present study were 1µg/ml & 1.5µg/ml and 0.094 µg/ml & 0.5 µg/ml respectively while in the Global Surveillance Program were 0.12 µg/ml& 4 µg/ml and 1 µg/ml & 2 µg/ml; and in the study conducted in Bangalore by Chandran et al 0.094 µg/ml & 0.125 µg/ml and 1 µg/ml & 2 µg/ml respectively [14,15]. This breakdown helps us identify, the MIC<sub>50</sub> of colistin among *Enterobacteriaceae* is high in the present study compared to the other studies indicating a impending rise of resistance in the near future, though the prevalence of colistin resistance is low currently. Thus, prompting implementation of appropriate approach in using these antibiotics.

In the present study, 100% and 95% of *Pseudomonas spp* and *Acinetobacter spp.* were susceptible to colistin with MIC<sub>50</sub> and MIC<sub>90</sub> being 0.75 µg/ml and 1.5 µg/ml respectively for both organism. While 70% of the *Acinetobacter spp.* were susceptible to Tigecycline with MIC<sub>50</sub> and MIC<sub>90</sub> being 0.094 µg/ml and 0.5µg/

ml. In a study conducted in Egypt 2015 by Wesam Hatem Amer, *Acinetobacter spp.* revealed 100% and 43.3% susceptibility to Colistin and Tigecycline respectively with MIC<sub>50</sub> and MIC<sub>90</sub> 0.5 µg/ml & 1 µg/ml and 2 µg/ml & 4 µg/ml respectively [16]. In the study conducted by Sangeetha et al in 2012 at Vellore revealed Colistin and Tigecycline MIC<sub>50</sub> and MIC<sub>90</sub> of *Pseudomonas spp.* and *Acinetobacter spp.* as 1 µg/ml & 2 µg/ml and 0.5 µg/ml & 64 µg/ml respectively [17]. These values strengthen the finding that the prevalence of resistance among NFGNB against colistin is much lower in the present study, though the same cannot be said for tigecycline but with proper guidelines the development of resistance can be delayed significantly.

There are few limitations in the study. There are chances for inaccuracies due to lack in use of genotypic methods. Polymyxins diffuses poorly in the media, hence the use of E-test tends to underestimate the resistance pattern.

## Conclusion

With the findings in the current study, it is concluded that colistin and Tigecycline can be considered as an option in treating infections with carbapenemase producing Gram negative bacteria, however national and global surveillances studies are recommended to regulate the treatment protocol and dose adjustment. Emerging resistance against these drugs are also to be kept in mind, as it leaves us with very minimal options to treat them. Hence their judicious use and regular monitoring of their susceptibility pattern are also recommended.

## References

1. World Health Organization (2017) Global priority list of antibiotic-resistant bacteria to guide research, discovery, and development of new antibiotics.
2. Nordmann P, Poirel L (2019) Epidemiology and Diagnostics of Carbapenem Resistance in Gram-negative Bacteria. Clin Infect Dis 69: S521-S528.
3. Chang YT, Lin CY, Chen YH, Hsueh PR (2015) Update on infections caused by *Stenotrophomonas maltophilia* with particular attention to resistance mechanisms and therapeutic options. Front Microbiol 6: 893.
4. Rodríguez-Martínez JM, Poirel L, Nordmann P (2009) Molecular epidemiology and mechanisms of carbapenem resistance in *Pseudomonas aeruginosa*. Antimicrob Agents Chemother 53: 4783-4788.
5. Meletis G (2016) Carbapenem resistance: overview of the problem and future perspectives. Ther Adv Infect Dis 3: 15-21.
6. Sekyere JO, Govinden U, Essack S (2016) The Molecular Epidemiology and Genetic Environment of Carbapenemases Detected in Africa. Microb Drug Resist 22: 59-68.
7. Nordmann P, Poirel L (2014) The difficult-to-control spread of carbapenemase producers among Enterobacteriaceae worldwide. Clin Microbiol Infect 20: 821-830.
8. Tascini C, Tagliaferri E, Giani T, Leonildi A, Flammini S, et al. (2013) Synergistic activity of colistin plus rifampin against colistin-resistant KPC-producing *Klebsiella pneumoniae*. Antimicrob Agents Chemother 57: 3990-3993.
9. Stein C, Makarewicz O, Bohnert JA, Pfeifer Y, Kesselmeier M, Hagel S, et al. (2015) Three Dimensional Checkerboard Synergy Analysis of Colistin, Meropenem, Tigecycline against Multidrug-Resistant Clinical *Klebsiella pneumoniae* Isolates. PLoS ONE 10: e0126479.
10. Munoz-Price LS, Poirel L, Bonomo RA, Schwaber MJ, Daikos GL, et al. (2013) Clinical epidemiology of the global expansion of *Klebsiella pneumoniae* carbapenemases. Lancet Infect Dis 13: 785-796.
11. EUCAST (2017) European Committee on Antimicrobial Susceptibility Testing breakpoint tables for interpretation of MICs and zone diameters, version 7.1.
12. BSAC (2016) BSAC methods for antimicrobial susceptibility testing.
13. Bialvaei AZ, Samadi Kafil H (2015) Colistin, mechanisms and prevalence of resistance. Curr Med Res Opin 31: 707-721.
14. Bradford PA, Kazmierczak KM, Biedenbach DJ, Wise MG, Hackel M, et al. (2015) Correlation of β-Lactamase Production and Colistin Resistance among *Enterobacteriaceae* Isolates from a Global Surveillance Program. Antimicrob Agents Chemother 60: 1385-1392.
15. Chandran S P, Nagaraj S, Kalal B S, Muralidharan S, Macaden R (2013) In-vitro susceptibility to colistin and tigecycline in New Delhi metallo-beta-lactamase-1 producing *Enterobacteriaceae*. Indian J Med Microbiol 31: 419-20.
16. Amer WH (2015) What remains against carbapenem-resistant *Enterobacteriaceae* and *Acinetobacter spp.*? Evaluation of Tigecycline and Colistin. Int J Curr Microbiol App Sci 4: 613-624.
17. Rajenderan S, Balaji V, Anandan S, Sahni RD, Tansarli GS, et al. (2014) Determination of MIC Distribution of Arbekacin, Cefminox, Fosfomicin, Biapenem and Other Antibiotics against Gram-Negative Clinical Isolates in South India: A Prospective Study. van Schaik W ed PLoS ONE 9: e103253.