Content available at: https://www.ipinnovative.com/open-access-journals



IP International Journal of Medical Microbiology and Tropical Diseases

Journal homepage: https://www.ijmmtd.org/

Original Research Article

Prevalence of multidrug-resistant *Acinetobacter baumannii* in endotracheal aspirate samples: Experience at a tertiary hospital

Dharnish Kumar Jha^{1,*}, Basudha Khanal², Ratna Baral²

¹Dept. of Microbiology, Koshi Hospital, Biratnagar, Nepal
 ²Dept. of Microbiology, B. P. Koirala Institute of Health Sciences, Dharan, Nepal



E PUBL

ARTICLE INFO

Article history: Received 06-05-2023 Accepted 01-06-2023 Available online 18-07-2023

Keywords: Intensive care unit Multidrug resistant Endotracheal aspirate Possible Ventilator associated pneumonia

ABSTRACT

Introduction: The emergence of resistance to multiple antimicrobial agents in pathogenic bacteria poses a significant public health threat because few or no effective antimicrobials are available for infectious diseases. *Acinetobacter baumannii* is a major cause of device-associated infections that pose a serious threat to critically ill patients. Resistance patterns are thought to result in very limited treatment options and high mortality. We examined the prevalence of *Acinetobacter baumannii* in endotracheal aspirates samples and explored their antibiotics susceptibility.

Objective: To determine the value of routine endotracheal aspirate cultures performed prior to the onset of the likely onset of ventilator-associated pneumonia (PVAP) in predicting pathogenic microorganisms and susceptibility to their antibiotics.

Materials and Methods: Patients admitted to the ventilatory intensive care unit were tested daily, and endotracheal aspirated (ET) specimens from suspected patients were sent to a microbiology laboratory for culture and sensitivity measurements.

Results: Of the 52 patients, only twenty five (48%) developed PVAP. Endotracheal aspirate cultures were positive in all PVAP cases. The most commonly isolated bacteria was *Acinetobacter baumannii* 14 (56%), followed by *Pseudomonas aeruginosa* 6 (24%) and *Klebsiella pneumoniae* 4 (16%). Almost all isolates of *Acinetobacter baumannii* are multidrug resistant (MDR). ICU stays greater than 16 days were observed for the pathogen *Acinetobacter baumannii*.

Conclusion: We believe that multidrug-resistant *Acinetobacter baumannii* is a widespread epidemic, leading to high mortality, long ICU stays, and a difficult case for ICU physicians. Further prospective studies are needed to tackle this threat.

This is an Open Access (OA) journal, and articles are distributed under the terms of the Creative Commons Attribution-NonCommercial-ShareAlike 4.0 License, which allows others to remix, tweak, and build upon the work non-commercially, as long as appropriate credit is given and the new creations are licensed under the identical terms.

For reprints contact: reprint@ipinnovative.com

1. Introduction

Acinetobacter baumannii (A. baumannii) is a Gramnegative, non-fermentative bacterium and an important opportunistic pathogen in the hospital setting. A. baumannii can cause a wide range of serious nosocomial infections, including ventilator-associated pneumonia, bloodstream infections, skin and soft tissue infections, wound infections, and urinary tract infections.¹ Hard to eradicate. this is A's fault *A. baumannii* has the ability to form strong biofilms on both biotic and abiotic surfaces.^{2–4}*A. baumannii* can survive for 3 days-5 months on dry, inanimate objectsand is hydrophobic to help bind to foreign materials such as plastics used in endovascular devices, catheters and ventilators.^{5–7} *A. baumannii* are resistant to multiple antimicrobial agents, including carbapenems, and multidrug resistance (MDR) is very common.⁸ Recently, they have become mostly drug-resistant (XDR) and pan-drug Resistance (PDR). Virtually drug-resistant *A. baumannii*

https://doi.org/10.18231/j.ijmmtd.2023.015 2581-4753/© 2023 Innovative Publication, All rights reserved.

^{*} Corresponding author. E-mail address: dharnishjha@gmail.com (D. K. Jha).

isolates are increasing rapidly.⁹ As a result, the World Health Organization (WHO) has declared that *A. baumannii* poses a serious public health threat and new antibiotics are urgently needed and classified as an important priority pathogen. Therefore, this study was conducted to determine the prevalence of *A. baumannii* and its resistance pattern in endotracheal aspirate isolates from ventilator (MV) patients with suspected of developing PVAP in critically ill patients.

2. Materials and Methods

This prospective study was conducted from January 2020 to December 2020 at the Department of Microbiology, B P Koirala Institute of Health Sciences (BPKIHS), Dharan, Nepal. Ethical approval for this study was obtained from the Microbiology and Institutional Review Board of BPKIHS in Dharan. Patients admitted to the ICU with MV were evaluated daily using the ventilator-related event surveillance criteria established by the Centers for Disease Control and Prevention/ National Healthcare Safety Network in 2013. A four-quadrant method followed by a semi-quantitative culture method was used.¹⁰ Gramscreened endotracheal aspirates were plated on blood agar, MacConkey agar, cultured aerobically at 35 °C, and growth was recorded as no growth, +1, +2, +3, +4. Bacterial isolates were identified by colony morphology, Gram stain results, and the results of various biochemical assays using standard microbiological techniques.¹¹ Determining of Antibiotic Susceptibility using Muellar Hinton Agar culture plates according to Institutional Guidelines (CLSI-2019). Susceptibility to colistin was determined using the colistin broth disc (CBDE) elution method recommended by CLSI.12

2.1. Data analysis

Of the 52 patients who gotten MV for ≥ 2 days, as it were 25 met criteria for PVAP. Result of ET suctions and pooled information from all 25 patients sent for microbiology were entered into a database utilizing MS Exceed expectations 2007. SPSS adaptation 20 was utilized for measurable investigation. Categorical factors were compared utilizing the chi-square test. For cells with an anticipated number less than 5, Fisher's correct test was utilized. A t-test was utilized to compare the implies of two autonomous tests. P-values less than 0.05 were measurably critical.

3. Results

Endotracheal aspirate cultures resulted in bacterial growth in all cases of PVAP. A total of 25 (48%) patients developed PVAP. A total of 9 (36%) patients had earlyonset PVAP and 16 (64%) patients had late-onset PVAP (P<0.001). Acinetobacter baumannii is associated with both early and late PVAP, but is mainly associated with late onset. All isolated organisms were Gram-negative bacilli (GNB), most commonly *Acinetobacter baumannii* 14(56%) followed by *Pseudomonas aeruginosa* 6(24%) and Klebsiella pneumonia 4(16%). *Acinetobacter baumannii* was the major pathogen isolated accounting for 56% of culture positive cases. All of the strains were resistant to amikacin, ampicillin, ceftazidime, tobramycin, piperacillin and carbenicillin whereas all were susceptible to colistin.

Table 1: Microorganisms isolated from ETA cultures in patients with PVAP (n=25).

Organism	Number of patients (%)
Acinetobacter baumannii	14(56)
Pseudomonas aeruginosa	6(24)
Klebsiella pneumoniae	4(16)
Escherichia coli	1(4)
Total	25(100)

Table 2: Number of organism isolated from early and late onset of PVAP

Onset of PVAP	Isolated organism	Number
Early-onset PVAP	Acinetobacter baumannii	6
	Pseudomonas aeruginosa	2
	Escherichia coli	1
Late-onset PVAP	Acinetobacter baumannii	8
	Pseudomonas aeruginosa	4
	Klebsiella pneumoniae	4

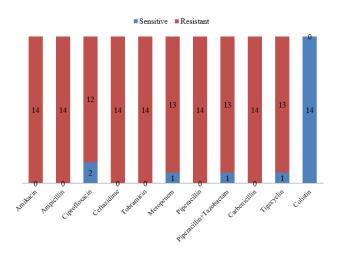


Fig. 1: Antibiotic susceptibility patterns of *Acinetobacter baumannii* to multiple antibiotics (n=14)

ICU length of stay ranged from 8 to 18 days, with a mean total length of stay of 11.2 days, although ICU length of stay was greater than 16 days observed for the pathogens *Acinetobacter baumannii* and *Pseudomonas aeruginosa*.

4. Discussion

Multidrug-resistant (MDR) pathogens are defined as resistant to three or more antibiotics.¹³ In addition, antibiotic-resistant infections are associated with longer hospital stays and higher medical costs than infections caused by antibiotic-sensitive strains (Cohen 1992). Polymyxin is the remaining antibiotic with relatively stable activity against MDR strains of Acinetobacter spp, P. aeruginosa, (McGowan 2006). All microorganisms isolated in our study were Gram-negative bacteria (GNB), Acinetobacter baumannii 14 (56%), Pseudomonas aeruginosa 6 (24%), Klebsiella pneumoniae 4 (16%) and Escherichia coli 1 (4%) in descending order in regularity. A study by John et al al.¹⁴ found that gram-negative bacteria were the most common pathogen associated with his PVAP in which, Acinetobacter species predominated (48.21%), followed by Klebsiella (19.64%). Pooling data from 24 published studies, Chastre and Fagon¹⁵⁻¹⁸ found that 58% of the isolates were Gram-negative bacteria, with the most common being Pseudomonas spp. followed by Acinetobacter and Proteus species. I found that Acinetobacter baumannii was the primary pathogen isolate accounting for 56% of culture-positive cases. All strains were multidrug-resistant. Similar results have been reported by John et al,¹⁴ The highest number of MDR isolates was Acinetobacter species (51.85%). A prospective study performed in a tertiary care hospital reported that Acinetobacter was the most common multidrug-resistant pathogen (47.9%), followed by Pseudomonas (27%).^{19,20} A study in India showed that 98.3% of A. baumannii causing ventilator-associated pneumonia was MDR-A. baumannii.²¹ Similarly, A 100% isolated A. baumannii was also the MDR in this study. In this study, the duration of ICU stay was > 16 days compared with A. baumannii infection and mostly associated with late-onset of PVAP. Similarly, in one study, the mean length of hospital stay for A. baumannii was approximately 24 days.²² In another study, the mean hospital stay of patients with A. baumannii is 20.25 days.²³ 23 In our study, a total of 25 (48%) patients had PVAP. A total of 9 patients (36%) had early-onset PVAP and 16 (64%) had late-onset PVAP (P < 0.001). A study conducted in India showed a total of 74 (27.71%) patients with PVAP. Of which, 13 (17.56%) patients had earlyonset VAP and 61 (82.43%) had late-onset VAP.²⁴In another study from India, 44.23% were classified as early VAP and 55.77% as late VAP, which is close to our finding.²⁵

5. Conclusion

In summary, the results of this study are: PVAP caused by MDR *A. baumannii* may be associated with longer hospital stays and intensive care days. Taken together, these results suggest that it is important to learn more about the behavior of MDR *A. baumannii* and try to prevent its spread. A.baumannii has become a hospital pathogen. The treatment options available for MDR *A. baumannii* infection are limited and colistin is often reported as the only effective antibiotic, making the pathogen very difficult to treat. Despite many efforts to study the mechanism of antibiotic resistance and its epidemiology, to date our understanding of its pathology is still relatively limited. Infection control surveillance, preventive measures, rigorous antibiotic prescribing, antibiotic resistance monitoring programs, and antibiotic cycles are essential for infection control in humans. Infection Prevention and Control (IPC) is essential and of great value in preventing infection and controlling this pathogen.

6. Source of Funding

None.

7. Conflicts of Interest

None.

Acknowledgment

We would like to sincerely thank all the staff of the Department of Microbiology of BPKIHS, Dharan for their assistance in conducting the study.

References

- 1. Moubareck CA, Halat DH. Insights into Acinetobacter baumannii: a review of microbiological, virulence, and resistance traits in a threatening nosocomial pathogen. *Antibiotics (Basel)*. 2020;9(3):119. doi:10.3390/antibiotics903011.
- 2. Lindford A, Kiuru V, Anttila VJ, Vuola J. Successful eradication of multidrug resistant Acinetobacter in the Helsinki Burn Centre. *J Burn Care Res.* 2015;36(6):595–601. doi:10.1097/BCR.00000000000209.
- 3. Gray AP, Allard R, Paré R, Tannenbaum T, Lefebvre B, Lévesque S, et al. Management of a hospital outbreak of extensively drug-resistant Acinetobacter baumannii using a multimodal intervention including daily chlorhexidine baths. *J Hospital Infect*. 2016;93(1):29–34.
- Eze EC, Chenia HY, and MEZ. Acinetobacter baumannii biofilms: effects of physicochemical factors, virulence, antibiotic resistance determinants, gene regulation, and future antimicrobial treatments. *Infect Drug Resist.* 2018;11:2277–99. doi:10.2147/IDR.S169894.
- Wieland K, Chhatwal P, Vonberg RP. Nosocomial outbreaks caused by Acinetobacter baumannii and Pseudomonas aeruginosa: Results of a systematic review. *Am J Infect Control.* 2018;46(6):643–8.
- Tacconelli E, Cataldo MA, Dancer SJ, De Angelis G, Falcone M, Frank U, et al. ESCMID guidelines for the management of the infection control measures to reduce transmission of multidrug-resistant Gram-negative bacteria in hospitalized patients. *Clin Microbiol Infect*. 2014;20(Suppl 1):1–55. doi:10.1111/1469-0691.12427.
- Asif M, Alvi IA, Rehman SU. Insight into Acinetobacter baumannii: pathogenesis, global resistance, mechanisms of resistance, treatment options, and alternative modalities. *Infect Drug Resist.* 2018;11:1249– 60. doi:10.2147/IDR.S166750.
- Nasr P. Genetics, epidemiology, and clinical manifestations of multidrug-resistant Acinetobacter baumannii. J Hosp Infect. 2020;104:4–11.
- 9. Teerawattanapong N, Panich P, Kulpokin D, Ranong SN, Kongpakwattana K, Saksinanon A, et al. A systematic review

of the burden of multidrugresistant healthcare-associated infections among intensive care unit patients in Southeast Asia: the rise of multidrug-resistant Acinetobacter baumannii. *Infect Control Hosp Epidemiol.* 2018;39(5):525–33. doi:10.1017/ice.2018.58.

- Bergmans DC, Borten MJ, De Lecuw P, Stobberingh EE. Reproducibility of quantitative cultures of endotracheal aspirates from mechanically ventilated patients. *J Clin Microbiol*. 1997;35(3):796–8. doi:10.1128/jcm.35.3.796-798.1997.
- Clinical Laboratory Standards Institute. Performance Standards for Antimicrobial Susceptibility Testing. 29th Edition. CLSI supplement M100. Wayne, PA: Clinical Laboratory Standards Institute; 2019.
- Humphries RM, Green DA, Schuetz AN, Bergman Y, Lewis S, Yee R, et al. Multicenter evaluation of colistin broth disk elution and colistin agar test: a report from the Clinical and Laboratory Standards Institute. *J Clin Microbiol.* 2019;57(11):e01269–19. doi:10.1128/JCM.01269-19.
- Magiorakos AP, Srinivasan A, Carey RB, Carmeli Y, Falagas ME, Giske CG, et al. Multidrug-resistant, extensively drug-resistant and pandrug-resistant bacteria: an international expert proposal for interim standard definitions for acquired resistance. *Clin Microbiol Infect*. 2012;18(3):268–81. doi:10.1111/j.1469-0691.2011.03570.x.
- Kapaganty VC, Pilli R. Microbiological profile of ventilatorassociated pneumonia in the intensive care unit of a tertiary hospital in Visakhapatnam, India. *Indian J Microbiol Res.* 2018;15(2):252–7. doi:10.18231/2394-5478.2018.0053.
- Chastre J, Fagon JY. Ventilator-associated pneumonia. *Am J Respir Crit Care Med.* 2002;165(7):867–903. doi:10.1164/ajrccm.165.7.2105078.
- Dey A, Bairy I. Incidence of multidrug-resistant organisms causing ventilator-associated pneumonia in a tertiary care hospital: A nine months' prospective study. *Ann Thorac Med.* 2007;2(2):52–7. doi:10.4103/1817-1737.32230.
- Baraibar J, Correa H, Mariscal D, Gallego M, Valles J, Rello J, et al. Risk factors for infection by Acinetobacter baumannii in intubated patients with nosocomial pneumonia. *Chest.* 1997;112(2):1050–4. doi:10.1378/chest.112.4.1050.
- Trouillet JL, Chastre J, Vuagnat A, Joly-Guillou ML, Combaux D, Dombret MC, et al. Ventilator-associated pneumonia caused by potentially drugresistant bacteria. *Am J Respir Crit Care Med.* 1998;157(2):531–9. doi:10.1164/ajrccm.157.2.9705064.
- Husni RN, Goldstein LS, Arroliga AC, Hall GS, Fatica C, Stoller JK, et al. Risk factors for an outbreak of multi-drug-resistant acinetobacter nosocomial pneumonia among intubated patients. *Chest.* 1999;115(5):1378–82. doi:10.1378/chest.115.5.1378.

- Garnacho-Montero J, Leyba CO, Fernandez-Hinojosa E, Aldaboallas T, Cayuela A, Marquez-Vacaro JA, et al. Acinetobacter baumannii ventilator-associated pneumonia: Epidemiological and clinical findings. *Intensive Care Med.* 2005;31(5):649–55. doi:10.1007/s00134-005-2598-0.
- Almomani B, Al-Gharaibeh R, Al-Mahasneh F, and SS. Multidrug resistant acinetobacter baumannii in ventilator associated pneumonia: Prevalence and predictors of mortality. *Eur Respir J.* 2014;44:2071.
- Perez F, Endimiani A, Ray AJ, Decker BK, Wallace CJ, Hujer KM, et al. Carbapenem-resistant Acinetobacter baumannii and Klebsiella pneumoniae across a hospital system: impact of post-acute care facilities on dissemination. *J Clin Microbiol*. 2010;65(8):1807–18. doi:10.1093/jac/dkq191.
- Alotaibi T, Abuhaimed A, Alshahrani M, Albdelhady A, Almubarak Y, Almasari O, et al. Prevalence of multidrug-resistant Acinetobacter baumannii in a critical care setting: a tertiary teaching hospital experience. SAGE Open Med. 2021;9:20503121211001144. doi:10.1177/20503121211001144.
- Patil HV, Patil VC. Incidence, bacteriology, and clinical outcome of ventilator-associated pneumonia at tertiary care hospital. *J Nat Sci Biol Med.* 2017;8(1):46–55. doi:10.4103/0976-9668.198360.
- Golia S, Sangeetha KT, Vasudha CL. Microbial profile of early and late onset ventilator associated pneumonia in the intensive care unit of a tertiary care hospital in bangalore, India. J Clin Diagn Res. 2013;7(11):2462–6. doi:10.7860/JCDR/2013/6344.3580.

Author biography

 Dharnish
 Kumar
 Jha,
 Consultant
 Microbiologist

 10
 https://orcid.org/0000-0002-3112-4829
 Microbiologist
 Microbiologist

Basudha Khanal, Professor D https://orcid.org/0000-0003-4340-6062

Ratna Baral, Additional Professor

Cite this article: Jha DK, Khanal B, Baral R. Prevalence of multidrug-resistant *Acinetobacter baumannii* in endotracheal aspirate samples: Experience at a tertiary hospital. *IP Int J Med Microbiol Trop Dis* 2023;9(2):77-80.