



Detection of beta-lactamase genes (bla_{TEM} and bla_{CTX}) glutaraldehyde in samples of *Acinetobacter baumannii*

Mojtaba Sade*1, Hussein Godarzi², Gita Eslami², Masoumeh Hallaj Zade², Fatemeh Fallah², Davood Yadegarnia³

1 International Branch Shahid Beheshti University of Medical Sciences and health services, Tehran, Iran

2 Microbiology group of the Shahid Beheshti University of Medical Sciences

3 Infectious Diseases and Tropical Medicine Research Center, Shahid Beheshti University of Medical Sciences, Tehran, Iran



Background and aims:

Excessive consumption of antimicrobial materials in hospitals is as the main encoder, led to the emergence, development and acquisition of new bacterial resistance to beta-lactamase. According to the lack of the enough information about the mechanism of the resistant genes to disinfectants in the country about this study and with the aim of considering the resistance or sensitivity of the isolates of the *Acinetobacter baumannii* (MDR) in facing 2% glutaraldehyde, this study was conducted in the selected intensive care units of the hospitals of Tehran in 2013.

Materials and Methods:

This study which was conducted over a period of 10 months, *Acinetobacter baumannii* species were separated by culture and biochemical tests from intensive care units of some hospitals in Tehran (Fayazbaksh, Taleghani, Imam Khomeini, Valiasr, Labafinejad). The resistance and sensitivity of the isolates to antibiotics is considered according to CLSI (2012) guidelines. By multiplex PCR method bla_{CTX} and bla_{TEM} were detected and finally, MDR strains were treated with 2% glutaraldehyde. PCR was put for each strains of MDR using specific primers.

Results:

In our study 131 isolates out of 588 (22/3%) of *Acinetobacter baumannii* were isolated. The amount of the resistance to various antibiotics was in the range of the 69/4% to 100%. The percentage of frequency of the bla_{TEM} and bla_{CTX} was 3/2% and 19/4% respectively. MIC_{50%} and MIC_{90%} of imipenem and meropenem antibiotics were $32 \pm 1 \mu\text{g}/\text{ml}$ and $64 \pm 1 \mu\text{g}/\text{ml}$ respectively ($P < 0.9$). And there was seen no resistance to glutaraldehyde. Some different band electrophoresis had been seen in the PCR of MDR strains.

Conclusion:

It seems that beside variety and prevalence of bla_{TEM} and bla_{CTX} , enormous mechanisms like porin and leaking systems (efflux Pumps) are responsible in the making of the resistance of *Acinetobacter baumannii* to disinfectants. The study about an accurate consideration of the resistance in strains and other microorganisms is advised.

Keywords: *Acinetobacter baumannii*, lactamase genes, disinfectants, antimicrobial resistance

References:

-Ferreira AE, Marchetti DP, Cunha GR, Oliveira LM, Fuentefria DB, Dall Bello AG, et al. Molecular characterization of clinical multiresistant isolates of *Acinetobacter* sp. From hospitals in Porto Alegre, State of Rio Grande do Sul, Brazil. *Rev Soc Bras Med Trop* 2011;44(6):725-30.

-Peymani A K, et al. Prevalence of class 1 integron among multidrug-resistant *Acinetobacter baumannii* in Tabriz, northwest of Iran. *Pol J Microbiol*. 2012;61(1):57-60.

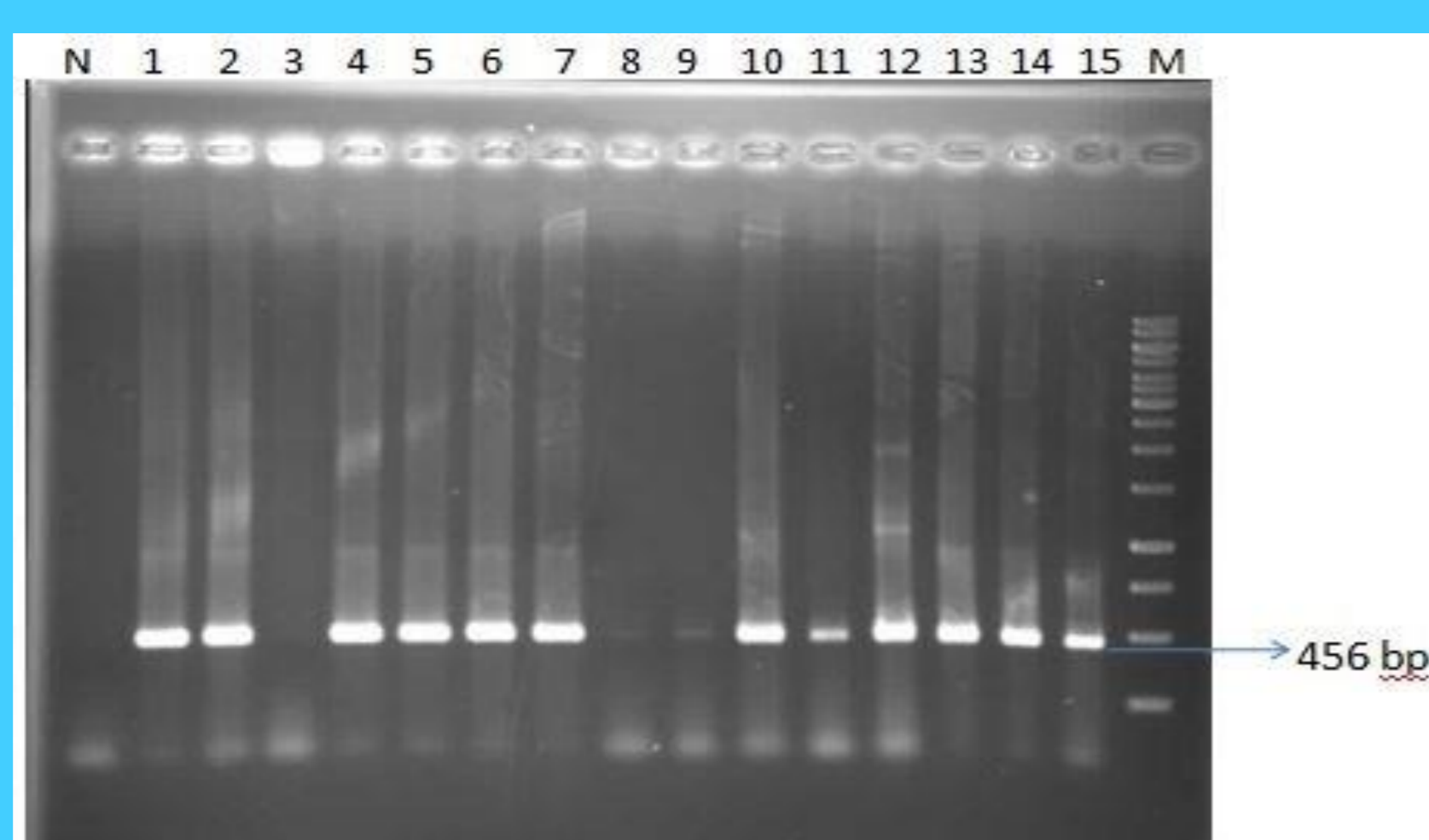


Figure 1: gel electrophoresis of *Acinetobacter baumannii* gene (bla_{CTX}) strains isolated from Surfaces and equipment N: Number of template DNA, M: size marker 1 kb ladder.

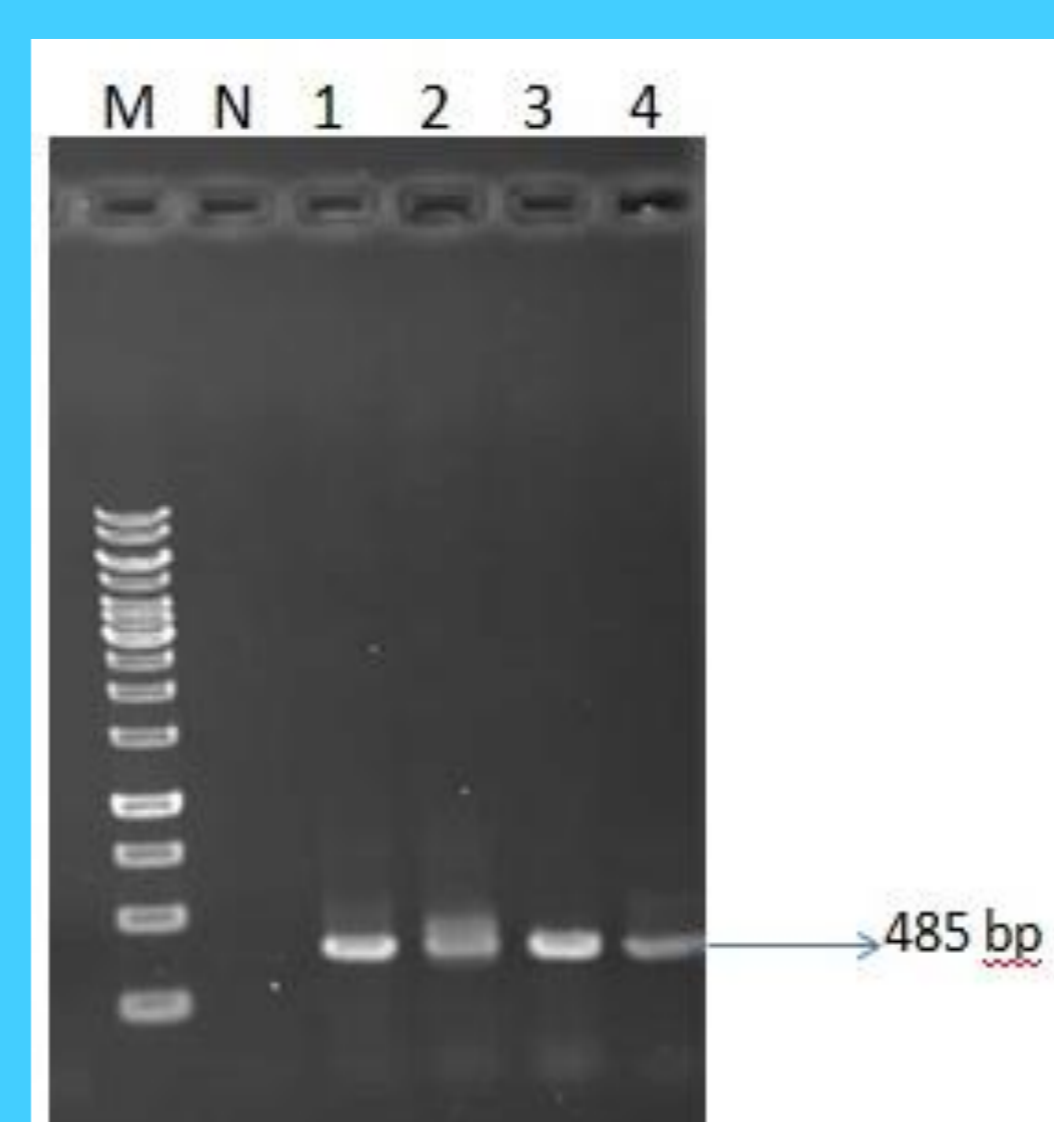


Figure 2: gel electrophoresis gene bla_{CTX} , M: size marker 1 kb ladder