The Lebanese LSIDCM

Combating antimicrobial resistance using antimicrobial combination therapy and β -lactamase inhibitors

Bassam El-Hafi^{1,2}, Sari Shawki Rasheed^{1,2}, Noor Salloum^{1,2}, Antoine Abou Fayad^{1,2}, George Farah Araj^{2,3}, Ghassan Matar Matar^{1,2}

¹ Department of Experimental Pathology, Immunology and Microbiology, American University of Beirut, Beirut, Lebanon

² Center for Infectious Disease Research (CIDR), American University of Beirut Medical Center, Beirut, Lebanon

³ Department of Pathology and Laboratory Medicine, American University of Beirut Medical Center, Beirut, Lebanon

Abstract

Introduction: The range of antimicrobial agents used to treat bacterial infections is becoming limited with the constant increase in antimicrobial resistance (AMR). Several genetic factors underlie AMR, including β -lactamase-encoding genes such as $bla_{CTXM-15}$ that confers resistance to third-generation cephalosporins, and bla_{OXA-48} , bla_{NDM-1} , and bla_{KPC-2} that confer resistance to carbapenems. Remaining treatment approaches for such resistant infections include antimicrobial combination therapy and the use of β -lactamase inhibitors. This study assesses the molecular effects of such treatment approaches on antimicrobial resistant *Enterobacteriaceae* clinical isolates *in vitro* and *in vivo*.

Methodology: Nine clinical *Enterobacteriaceae* isolates were included in the study. One harboring $bla_{CTXM-15}$, one harboring bla_{OXA-48} , one harboring bla_{KPC-2} , two harboring bla_{NDM-1} and $bla_{CTXM-15}$, and four harboring bla_{OXA-48} and $bla_{CTXM-15}$. Minimal inhibitory concentrations were determined for carbapenems with β -lactamase inhibitors: avibactam, Ca-EDTA, and relebactam. Synergism between antibiotic combinations was determined by double disc diffusion when using colistin with several antibiotics. *In vitro* and *in vivo* gene expression levels were done on these combinations with and without inhibitors.

Results: The use of meropenem, imipenem, and ertapenem with the selected β -lactamase inhibitors restored isolate susceptibility in 100%, 87.5%, and 25% of the cases, respectively. Antimicrobial synergism was mostly detected between colistin and meropenem, fosfomycin, or tigecycline. Survival studies revealed the survival of most mice receiving antimicrobial combination therapy with inhibitors as compared to the controls. Overall gene expression levels of resistance genes were variable depending on treatment.

Conclusions: The threat of antibiotic resistant bacterial infections remains viable; however, different approaches to therapy are available.

Key words: antimicrobial resistance; combination therapy; beta-lactamase inhibitor.

J Infect Dev Ctries 2018; 12(2S):14S. doi:10.3855/jidc.10099

(Received 21 December 2017 – Accepted 22 December 2017)

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Corresponding author

Ghassan M. Matar, M.S., Ph.D. Department of Experimental Pathology, Immunology and Microbiology Laboratory Director, Center for Infectious Diseases Research (CIDR) American University of Beirut Riad El-Solh St. P.O.BOX 11-0236 Beirut 1107 2020 Lebanon Phoone: +961 1 350 000 Ext. 5128 E-mail: gmatar@aub.edu.lb

Conflict of interests: No conflict of interests is declared.