

## Acquisition of Colistin Resistance Links Cell Membrane Thickness Alteration with a Point Mutation in Lpxd Gene in Acinetobacter BaumanniiLinn

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## Abstract

Acinetobacter baumannii is one of the most common causes of nosocomial infections in intensive

care units. Its ability to acquire diverse mechanisms of resistance limits its therapeutic choices. Multi-drug resistant A. baumannii became a critical challenge for common antibiotic treatments. Colistin is an old antibiotic that has been reused recently, as a last resort drug, for A. baumannii. In this study, we explored the impact of gaining colistin resistance on the susceptibility to other antibiotics and linking the resistance acquisition to its genetic basis. The susceptibility of 95 A. baumannii isolates revealed that 89 isolates were MDR with five isolates resistant to colistin. Subsequently, three isolates, -MS48, MS50, and MS64-, exhibited different resistance pattern, when exposed to colistin resistance induction. The susceptibility profiles reevaluation showed the gain of resistance to almost all tested antibiotics for MS50Col-R and MS64Col-R isolates, while the resistance pattern of MS48Col-R remained unchanged. Upon TEM examination, morphological alterations compared to parent sensitive strains, have been reported for all of the induced isolates, also to an isolated colistin-resistant clinical isolate (MS34Col-R). Finally, genetic alterations in PmrB and LpxACD were assessed, in which a point mutation in LpxD in MS64Col-R mutant and the colistin-resistant isolate MS34Col-R at Lys117Glu was identified in the lipid-binding domain. Our findings shed light on the implications of using colistin in the treatment of A. baumannii, especially in sub-MIC concentrations, where co-resistance to other classes of antibiotics may emerge beside the rapid acquisition of resistance against colistin itself due to distinct genetic events



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