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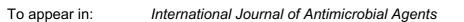
Diverse genetic alterations responsible for post-exposure colistin resistance in populations of the same strain of Klebsiella pneumoniae

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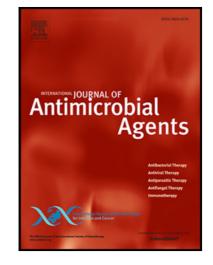
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Highlights

• We investigated the intra- and inter-strain diversity of genetic alterations in *Klebsiella pneumoniae* after exposure to colistin.

• We found diverse amino acid alterations associated with colistin resistance between the *K*. *pneumoniae* strains.

• The mutations varied between single-step colistin-resistant mutants derived from the same strain.

• We found colistin-resistant mutants with no amino acid changes in the PmrAB or PhoPQ two-component regulatory systems, and while some mutants showed *pbgP* mRNA overexpression, others did not.

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