

Genomic analysis unveils important aspects of population structure, virulence, and antimicrobial resistance in *Klebsiella aerogenes*

Hemanoel Passarelli Araujo, Jussara Kasuko Palmeiro, Kanhu Charan Moharana, Francisnei Pedrosa da Silva, Libera Maria Dalla Costa, Thiago Venancio

Abstract

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