

Review of: "Mobilome-driven segregation of the resistome in biological wastewater treatment"

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This article provides a comprehensive and meta-omic analysis of sequence data from a Biological wastewater treatment plant, with focus on plasmids and phages as the major mobile genetic elements involved in the dissemination of various antibiotic resistance genes (ARGs). The manuscript is well written with few minor grammatical errors, the only limitation to the manuscript is the lack of specific details of the ARGs identified corresponding to the major categories of antimicrobial classes described. There is also no specific details of the types of plasmids (incompatibility groups) and phages described in the paper. I believe this missing information is critical, and will increase the impact of the data that is described.