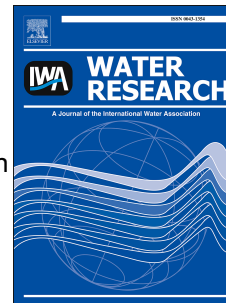


# Accepted Manuscript

Metagenomic binning reveals the functional roles of core abundant microorganisms in twelve full-scale biogas plants

Stefano Campanaro, Laura Treu, Panagiotis G. Kougias, Gang Luo, Irini Angelidaki



PII: S0043-1354(18)30337-3

DOI: [10.1016/j.watres.2018.04.043](https://doi.org/10.1016/j.watres.2018.04.043)

Reference: WR 13740

To appear in: *Water Research*

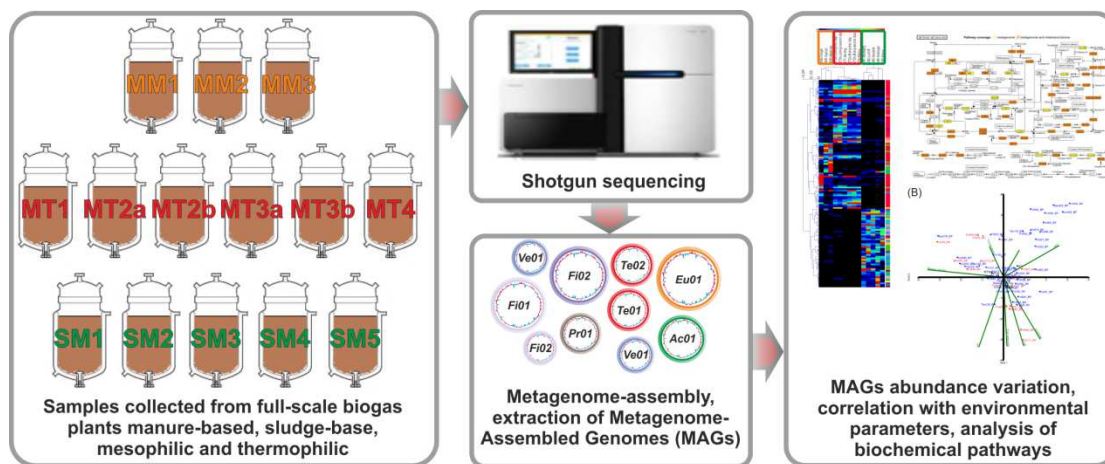
Received Date: 13 January 2018

Revised Date: 22 March 2018

Accepted Date: 16 April 2018

Please cite this article as: Campanaro, S., Treu, L., Kougias, P.G., Luo, G., Angelidaki, I., Metagenomic binning reveals the functional roles of core abundant microorganisms in twelve full-scale biogas plants, *Water Research* (2018), doi: 10.1016/j.watres.2018.04.043.

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.



Abstract Graphic

Download English Version:

<https://daneshyari.com/en/article/8873826>

Download Persian Version:

<https://daneshyari.com/article/8873826>

[Daneshyari.com](https://daneshyari.com)