

CORRECTION

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Correction: Thermochemical wastewater valorization *via* enhanced microbial toxicity tolerance

Lahiru N. Jayakody,^a Christopher W. Johnson,^a Jason M. Whitham,^b Richard J. Giannone,^b Brenna A. Black,^a Nicholas S. Cleveland,^a Dawn M. Klingeman,^b William E. Michener,^a Jessica L. Olstad,^a Derek R. Vardon,^a Robert C. Brown,^c Steven D. Brown,^{bd} Robert L. Hettich,^b Adam M. Guss^b and Gregg T. Beckham^{ab*}

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Correction for 'Thermochemical wastewater valorization *via* enhanced microbial toxicity tolerance' by Lahiru N. Jayakody *et al.*, *Energy Environ. Sci.*, 2018, 11, 1625–1638, DOI: 10.1039/C8EE00460A.

In the original version of the manuscript, the intergenic region between PP_1584 and PP_1585 was indicated in Table S8 as the integration site for *groES*, *groEL*, and *clpB* into the genome of strain LJ15. However, these genes were integrated at a different location in the genome. The text in Table S8 lists "EM42 with the *clpB-groES-groEL* chaperone expression cassette integrated within the intergenic region between PP_1584 and PP_1585". This text should be corrected to "EM42 with the *clpB-groES-groEL* chaperone expression cassette integrated within the intergenic region between **PP_4217 and PP_4218**".

Accordingly, the primers listed for confirming integration in the main text Materials and methods section "Strain construction" should be changed. The original text states: "Following sucrose selection, single colonies were subjected to colony PCR with primers oLJ154 (Fwd) and oLJ155 (Rev) to identify those with genome integration of the chaperone genes." This text should be corrected to "Following sucrose selection, single colonies were subjected to colony PCR with primers oLJ154 (Fwd) and oLJ155 (Rev) for **LJ14 and oCJ311 (Fwd) and oCJ312 (Rev) for LJ15** to identify those with genome integration of the chaperone genes." The primer sequences of oCJ311 and oCJ312 are not listed in the ESI, and the following entries should have been included in Table S10.

Primer	Sequence [5'-3']
oCJ311	agcctcttcagcgtcaac
oCJ312	cacgctgcttcattgaac

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

^a National Bioenergy Center, National Renewable Energy Laboratory, Golden, CO 80401, USA. E-mail: gregg.beckham@nrel.gov

^b Oak Ridge National Laboratory, 1 Bethel Valley Rd, Oak Ridge, TN 37830, USA

^c Bioeconomy Institute and Department of Mechanical Engineering, Iowa State University, Ames, IA 50011, USA

^d LanzaTech, Inc., Skokie, IL 60077, USA

