

Correction to GeoChip-Based Analysis of Microbial Functional Gene Diversity in a Landfill Leachate-Contaminated Aquifer

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Supporting Information

A number of errors were discovered after the original article was published.

Text Corrections:

1. Page 5824, the last sentence in the first paragraph should read "Since most municipalities at some time had facilities for MSW disposal..."
2. The location of Shimadzu Corporation on page 5825 should be Kyoto, Japan.
3. On Page 5825 the comma between "Millipore" and "Corp." should be removed.
4. The second sentence in Section 3.1 on page 5826 should read "The entire flowpath was suboxic (DO < 0.30 mg/L)..." (< replaced with ≤).
5. "CH₄, NH₄⁺" in the third sentence in Section 3.1 should be "H₂S"
6. Table 1 on page 5826, parentheses should be around the units for EC.
7. Table 2 on page 5827, An additional footnote should be added: "The number in parentheses represents the percentage of the genes in total gene numbers"
8. Page 5827, Section 3.3, second sentence, "535" should be "674".
9. Page 5829, Section 3.6, "nagG" should be "nagI", "0.42" should be "0.32", "0.05" should be "0.1", and "CH₄ and" should be removed.
10. The top sentence on page 5831 should read "...for the wells closest to the slough" ("closest" replaced with "close" and the last sentence, please add 'the' before 'microarray approach' and 'stable isotope probing method'.
11. The last sentence on page 5831 should read "Hopefully, the results can lead to more hypothesis-driven studies, such as the detection of functional activity using metatranscriptomics or the microarray approach,^{56–58} and the detection of active microorganisms using the stable isotope probing (SIP) method.⁵⁹"

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The Supporting Information is also corrected.

ASSOCIATED CONTENT

Supporting Information

This material is available free of charge via the Internet at <http://pubs.acs.org>.

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