ERRATUM

Erratum to: A molecular phylogenetic framework for *Bacillus subtilis* using genome sequences and its application to *Bacillus subtilis* subspecies *stecoris* strain D7XPN1, an isolate from a commercial food-waste degrading bioreactor

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In the online published article, the *Bacillus subtilis* subsp. *stecoris* name was wrongly published and it should be corrected to *Bacillus subtilis* subsp. *stercoris* throughout the manuscript.

Description of *Bacillus subtilis* subsp. *stercoris* subsp. nov

Bacillus subtilis subsp. *stercoris* [ster'co.ris. L. gen. n. stercoris, of compost, from which the strain was isolated]. Grows optimally at 45 °C (range 24–50), pH of 7 (range 5–9), and grew in the presence of 7% NaCl. Facultative anaerobe by fermentation, forms white irregular colonies 1–2 mm in diameter when grown on dTSA, cells are straight rods 4–5 µm length by 1 µm width, and stain Gram positive. Degrades potato starch by amylase activity and expresses partial β-xylanase activity detected when grown on dTSA with x-β-D-xyloside. Type strain is D7XPN1^T (=KCTC 33554^T, JCM 30051^T).

The online version of the original article can be found under doi:[10.1007/s13205-016-0408-8](https://doi.org/10.1007/s13205-016-0408-8).

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