Correction to: Bacterial association observations in *Lucilia sericata* and *Lucilia cuprina* organs through 16S rRNA gene sequencing

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Published online: 9 March 2021

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Correction to: Applied Microbiology and Biotechnology (2021) 105:1091–1106
<https://doi.org/10.1007/s00253-020-11026-8>

The published online version contains mistake in Figures 1 and 2. The researcher's regret that a publishing error has occurred in this article. Figure 1 was omitted and therefore does not appear in text. Figure 2 was also published missing key x-axis information.

Figures 1 and 2 should appear as follows:

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The online version of the original article can be found at <https://doi.org/10.1007/s00253-020-11026-8>

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Fig. 1 Total OTU count of bacterial sequences identified across all preliminary samples. Patterns denote organ sampled from specimens

Fig. 2 Phyla diversity by sample through 16S rDNA sequencing: M = male, F = female, WM = Westmead, DT = Digestive tract, SG = Salivary gland, Ovar = Ovaries, Sperma = Spermatheca, Sem Ves = Seminal vesicle