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Correction: Specific detection of *Cronobacter sakazakii* in powdered infant formula using ssDNA aptamer

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Correction for 'Specific detection of *Cronobacter sakazakii* in powdered infant formula using ssDNA aptamer' by Hyeyoung Kim et al., *Analyst*, 2021, **146**, 3534–3542, DOI: 10.1039/D1AN00118C.

The authors regret that there were errors in the buffer compositions described in section 2.3 of the article.

On page 3535, the sentence beginning "Briefly, the DNA library pool..." should be correctly given as "Briefly, the DNA library pool was denatured in binding buffer (25 mM of glucose, 5 mM of MgCl₂, 1 mg mL⁻¹ of BSA, and 0.1 mg mL⁻¹ of tRNA in pre-autoclaved 1× phosphate-buffered saline (PBS)) at 95 °C for 5 min and immediately chilled in an ice-bath before allowing the binding reaction with the target bacteria."

The sentence beginning "After a final partitioning step..." should be correctly given as "After a final partitioning step, bound ssDNA was eluted from the target bacteria by incubating the sample with 500 µL of elution buffer (25 mM of glucose, 5 mM of MgCl₂, and 1 mg mL⁻¹ of BSA in pre-autoclaved 1× PBS buffer) at 95 °C for 10 min."

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



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