

## CORRECTION

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## Correction: Optimisation of the dibromomaleimide (DBM) platform for native antibody conjugation by accelerated post-conjugation hydrolysis

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Correction for 'Optimisation of the dibromomaleimide (DBM) platform for native antibody conjugation by accelerated post-conjugation hydrolysis' by Maurício Morais *et al.*, *Org. Biomol. Chem.*, 2017, **15**, 2947–2952, DOI: 10.1039/C7OB00220C.

Following publication of this article, the authors discovered the NMR spectra of 7 compounds presented in the original ESI were inappropriately modified by a co-author using NMR processing software to remove peaks corresponding to minor impurities.

In addition, a SDS-PAGE gel control lane, containing the commercially purchased antibody, had also been partially cut from another gel for aesthetics purposes.

An investigation by UCL has concluded that this was a case of a researcher cleaning up NMR spectra for publication, without the knowledge of the corresponding authors, and the raw data is accurate.

Replacement spectra reproduced from the original raw NMR FID files have been provided in a revised ESI file for:

<sup>1</sup>H NMR: **4, 5, 8, 13, 14, h5**

<sup>13</sup>C NMR: **4, 5, 6, 8**

The SDS-PAGE gel control lane in Fig. S2 is also replaced with the original image.

The authors confirm the validity of the replacement spectra/images in the revised ESI in comparison to the originally published spectra/images. While some of the compounds do contain impurities, the amount does not affect the main substance or conclusions of the paper.

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

