

CORRECTION

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rsc.li/medchemCorrection for ‘Minimising the payload solvent exposed hydrophobic surface area optimises the antibody–drug conjugate properties’ by Adrian D. Hobson *et al.*, *RSC Med. Chem.*, 2024, <https://doi.org/10.1039/d3md00540b>.

The authors regret that there is an error in the structure of compound 3 in Fig. 2 of their manuscript. The 3-aminobenzyl group should be in the *para* position instead of the *meta* position as is shown. The corrected scheme is shown below.

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

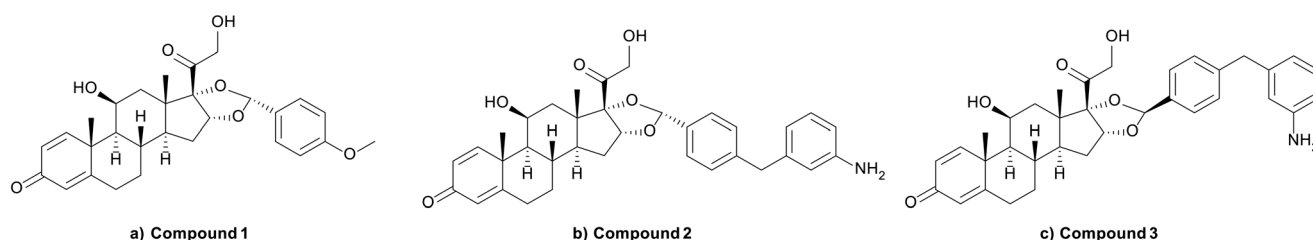


Fig. 2 GRM compounds a) mono aryl anisole compound 1, b) biaryl aniline with (*R*)-acetal compound 2, c) biaryl aniline with (*S*)-acetal compound 3.

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