

Author Correction: Second-generation DNA-templated macrocycle libraries for the discovery of bioactive small molecules

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Correction to: Nature Chemistry https://doi.org/10.1038/s41557-018-0033-8, published online 02 April 2018.

In the Supplementary Information file originally published for this Article there were several errors.

In the section 'Typical procedure of preparative split ligation', the sentence "30–40% yields" should have read "30–40% yields (w/ no extra ATP added)".

In the section 'Large-scale preparative splint ligation', the sentence "Likewise, the additional ATP should be added for higher yields" was missing from the end of the paragraph.

In the section 'Library assembly', the sentence "After 2 h at 30 °C, each batch was mixed with 6.5 mL (25% v/v) of 1 M NaOH solution, which was quenched with the same volume (6.5 mL) of 3 M NaOAc buffer pH 5.2" should have read "After 2 h at 30 °C, each batch was mixed with 6.5 mL (25% v/v) of 1 M NaOH solution and left for 30 minutes, followed by quenching with the same volume (6.5 mL) of 3 M NaOAc buffer pH 5.2." In the same section, the sentence "The solution in each conical was diluted just with 25 mL of isopropanol and passed through 14 HiBind DNA Midi columns (Omega Bio-tek)" should have read "The solution in each conical was diluted just with 25 mL of isopropanol and passed through 14 HiBind DNA Midi columns (7 columns per conical, Omega Bio-tek)."

In the section 'General procedure for selections and high-throughput sequencing', in the second paragraph, " \pm 5 mM DTT" should have read " \pm 2 mM TCEP" in two instances.

These errors have now been corrected.

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