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Correction: Aptamer-based strategies for recognizing adenine, adenosine, ATP and related compounds

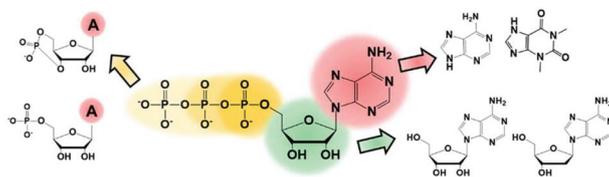
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Correction for 'Aptamer-based strategies for recognizing adenine, adenosine, ATP and related compounds' by Yuqing Li *et al.*, *Analyst*, 2020, **145**, 6753–6768, DOI: 10.1039/D0AN00886A.

The authors regret unfortunate errors in their paper on pages 6754 and 6760. The nucleoside structures are missing the oxygen in their sugar rings in Fig. 1, 2A, 8B and the graphical abstract. In addition, the L-enantiomer should be the D-enantiomer in Fig. 2A. The seven bases in the 3'-end of Fig. 2D are missing. The corrected Fig. 1, 2, 8B and graphical abstract figure are given below.



Graphical abstract figure

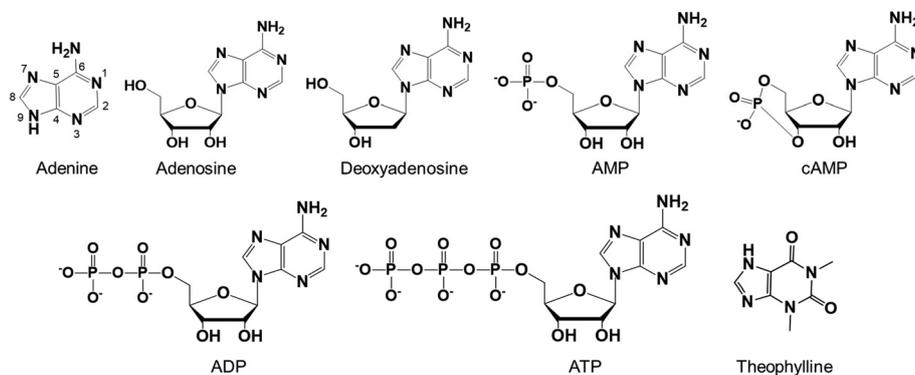


Fig. 1 Chemical structures of adenine and its derivatives reviewed including adenosine, deoxyadenosine, AMP, cAMP, ADP, ATP and theophylline.



