Correction to: The role of NMR in leveraging dynamics and entropy in drug design

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Unfortunately, in the original publication, Fig. 5 was published incorrectly. The correct version is given below:

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Fig. 5 An overview of the different components that contributes binding free energy in typical protein-ligand interaction and the relevant NMR experiments to study these individual contributions. It should

be noted that some of methods would not provide absolute measurements of these thermodynamic quantities, rather relative measurements across a panel of lead compounds and their analogs