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The authors regret that the structure of staurosporine is incorrect in Fig. 8 of the original article. A corrected version of Fig. 8, in which the endocyclic oxygen of the pyranose ring is now included, is presented herein.

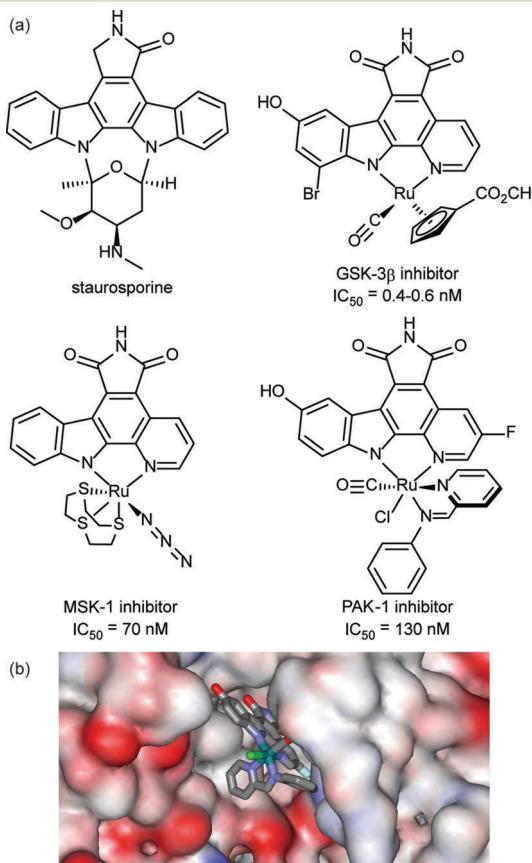


Fig. 8 Co-ordination complexes for kinase recognition (a) structures of different inhibitors highlighting similarity to natural product staurosporine – a pan kinase inhibitor (b) X-ray crystal structure of a ruthenium complex bound to PAK1 kinase domain (PDB ID: 3FXZ).¹⁰³

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

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