

# Correction to “Design and Function of Supramolecular Recognition Systems Based on Guest-Targeting Probe-Modified Cyclodextrin Receptors for ATP”

Kyohhei Fujita, Shoji Fujiwara, Tatsuru Yamada, Yuji Tsuchido,<sup>ⓑ</sup> Takeshi Hashimoto, and Takashi Hayashita\*<sup>ⓑ</sup>

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The ATP structure depicted in the Graphical Abstract and also in Figures 3 and 8 is incorrect; the structure is the enantiomer (mirror image) of natural ATP by mistake. Thus, the Graphical Abstract and Figures 3 and 8 should be corrected as follows:

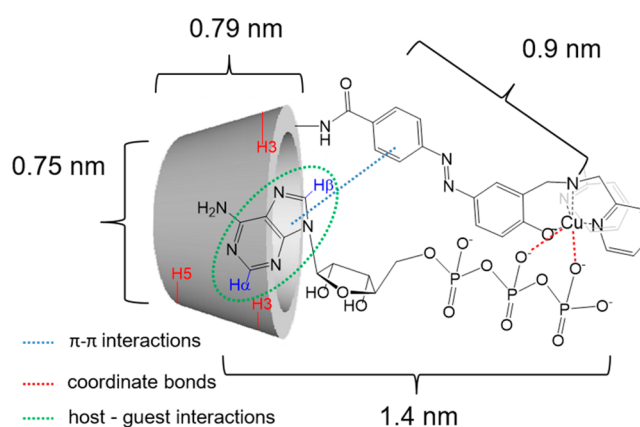
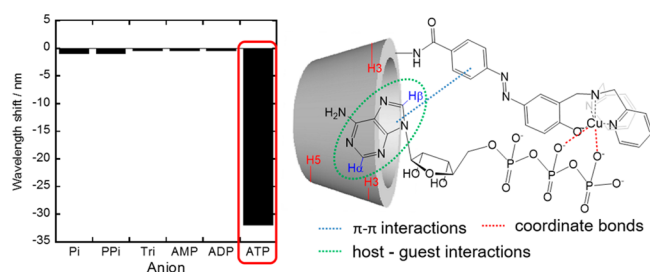


Figure 8.

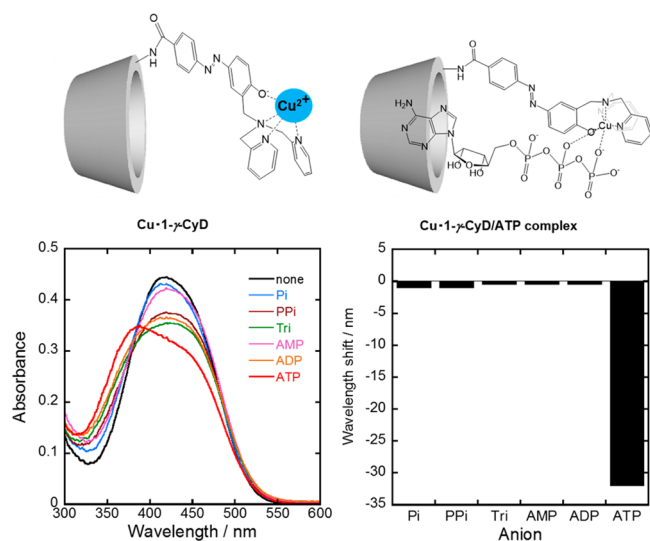


Figure 3.